



SEQUENCE LISTING

<110> Sylvain
Martin, Beauchamp
Quiniou, Christiane

<120> Cytokine receptors modulators, method of identifying same, and
method of modulating cytokine receptors activity with same

<130> GOUD:040US

<140> 10/693,657

<141> 2003-10-24

<150> 60/420,679

<151> 2002-10-24

<150> 60/423,530

<151> 2002-11-05

<160> 65

<170> PatentIn version 3.2

<210> 1

<211> 14

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 1

Gly Val Leu Ile Ile Ile Glu Leu Asn Thr Lys Glu Gln Ala
1 5 10

<210> 2

<211> 10

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 2

Glu Ala Thr Val Gly Glu Arg Val Arg Leu
1 5 10

<210> 3

<211> 10

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 3

Leu Pro Leu Glu Ser Asn His Thr Leu Lys
1 5 10

<210> 4

<211> 11

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 4

Ser Pro Val Asp Ser Tyr Gln Tyr Gly Thr Thr
1 5 10

<210> 5

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 5

Val Ile Leu Thr Asn Pro Ile Ser Lys Glu
1 5 10

<210> 6

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 6

Asn Lys Val Gly Arg Gly Glu Arg Val Ile
1 5 10

<210> 7
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<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 7

Met Pro Pro Thr Glu Gln Glu Ser Val
1 5

<210> 8
<211> 9
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 8

Arg Lys Thr Lys Lys Arg His Cys Val
1 5

<210> 9
<211> 9
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 9

Thr Val Leu Glu Arg Val Ala Pro Thr
1 5

<210> 10
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Peptide

<400> 10

Thr Ser Ile Gly Glu Ser Ile Glu Val

1

5

<210> 11
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 11

Ser Ile Phe Val Pro Arg Pro Glu Arg Lys
1 5 10

<210> 12
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 12

Asn Phe Leu His Asn Ser Ile Phe Val
1 5

<210> 13
<211> 9
<212> PRT
<213> Artificial Sequence

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Peptide
<400> 13

Glu Gly Pro Cys Pro Lys Val Cys Glu
1 5

<210> 14
<211> 10
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 14

Glu Ser Asp Val Leu His Phe Thr Ser Thr
1 5 10

<210> 15
<211> 9
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 15

Arg Thr Asn Ala Ser Val Pro Ser Ile
1 5

<210> 16
<211> 9
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<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 16

Ile Arg Lys Tyr Ala Asp Gly Thr Ile
1 5

<210> 17
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Peptide

<400> 17

Glu Asn Phe Ile His Leu Ile Ile Ala
1 5

<210> 18
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<213> Artificial Sequence

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Peptide

<400> 18

Ala Lys Thr Gly Tyr Glu Asn Phe Ile His
1 5 10

<210> 19

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 19

Lys Glu Arg Thr Val Ile Ser Asn Leu Arg
1 5 10

<210> 20

<211> 1356

<212> PRT

<213> Homo sapiens

<400> 20

Met Gln Ser Lys Val Leu Leu Ala Val Ala Leu Trp Leu Cys Val Glu
1 5 10 15

Thr Arg Ala Ala Ser Val Gly Leu Pro Ser Val Ser Leu Asp Leu Pro
20 25 30

Arg Leu Ser Ile Gln Lys Asp Ile Leu Thr Ile Lys Ala Asn Thr Thr
35 40 45

Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro
50 55 60

Asn Asn Gln Ser Gly Ser Glu Gln Arg Val Glu Val Thr Glu Cys Ser
65 70 75 80

Asp Gly Leu Phe Cys Lys Thr Leu Thr Ile Pro Lys Val Ile Gly Asn
85 90 95

Asp Thr Gly Ala Tyr Lys Cys Phe Tyr Arg Glu Thr Asp Leu Ala Ser
100 105 110

Val Ile Tyr Val Tyr Val Gln Asp Tyr Arg Ser Pro Phe Ile Ala Ser
115 120 125

Val Ser Asp Gln His Gly Val Val Tyr Ile Thr Glu Asn Lys Asn Lys
130 135 140

Thr Val Val Ile Pro Cys Leu Gly Ser Ile Ser Asn Leu Asn Val Ser
145 150 155 160

Leu Cys Ala Arg Tyr Pro Glu Lys Arg Phe Val Pro Asp Gly Asn Arg
165 170 175

Ile Ser Trp Asp Ser Lys Lys Gly Phe Thr Ile Pro Ser Tyr Met Ile
180 185 190

Ser Tyr Ala Gly Met Val Phe Cys Glu Ala Lys Ile Asn Asp Glu Ser
195 200 205

Tyr Gln Ser Ile Met Tyr Ile Val Val Val Val Gly Tyr Arg Ile Tyr
210 215 220

Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu
225 230 235 240

Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile
245 250 255

Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu
260 265 270

Val Asn Arg Asp Leu Lys Thr Gln Ser Gly Ser Glu Met Lys Lys Phe
275 280 285

Leu Ser Thr Leu Thr Ile Asp Gly Val Thr Arg Ser Asp Gln Gly Leu
290 295 300

Tyr Thr Cys Ala Ala Ser Ser Gly Leu Met Thr Lys Lys Asn Ser Thr
305 310 315 320

Phe Val Arg Val His Glu Lys Pro Phe Val Ala Phe Gly Ser Gly Met
325 330 335

Glu Ser Leu Val Glu Ala Thr Val Gly Glu Arg Val Arg Ile Pro Ala

340																345																350															
Lys	Tyr	Leu	Gly	Tyr	Pro	Pro	Pro	Glu	Ile	Lys	Trp	Tyr	Lys	Asn	Gly																																
		355						360					365																																		
Ile	Pro	Leu	Glu	Ser	Asn	His	Thr	Ile	Lys	Ala	Gly	His	Val	Leu	Thr																																
		370						375					380																																		
Ile	Met	Glu	Val	Ser	Glu	Arg	Asp	Thr	Gly	Asn	Tyr	Thr	Val	Ile	Leu																																
385								390					395																																		
Thr	Asn	Pro	Ile	Ser	Lys	Glu	Lys	Gln	Ser	His	Val	Val	Ser	Leu	Val																																
				405								410			415																																
Val	Tyr	Val	Pro	Pro	Gln	Ile	Gly	Glu	Lys	Ser	Leu	Ile	Ser	Pro	Val																																
				420					425					430																																	
Asp	Ser	Tyr	Gln	Tyr	Gly	Thr	Thr	Gln	Thr	Leu	Thr	Cys	Thr	Val	Tyr																																
				435					440					445																																	
Ala	Ile	Pro	Pro	Pro	His	His	Ile	His	Trp	Tyr	Trp	Gln	Leu	Glu	Glu																																
				450					455					460																																	
Glu	Cys	Ala	Asn	Glu	Pro	Ser	Gln	Ala	Val	Ser	Val	Thr	Asn	Pro	Tyr																																
465								470					475			480																															
Pro	Cys	Glu	Glu	Trp	Arg	Ser	Val	Glu	Asp	Phe	Gln	Gly	Gly	Asn	Lys																																
				485								490			495																																
Ile	Glu	Val	Asn	Lys	Asn	Gln	Phe	Ala	Leu	Ile	Glu	Gly	Lys	Asn	Lys																																
				500					505					510																																	
Thr	Val	Ser	Thr	Leu	Val	Ile	Gln	Ala	Ala	Asn	Val	Ser	Ala	Leu	Tyr																																
				515					520					525																																	
Lys	Cys	Glu	Ala	Val	Asn	Lys	Val	Gly	Arg	Gly	Glu	Arg	Val	Ile	Ser																																
				530								540																																			
Phe	His	Val	Thr	Arg	Gly	Pro	Glu	Ile	Thr	Leu	Gln	Pro	Asp	Met	Gln																																
545								550					555			560																															
Pro	Thr	Glu	Gln	Glu	Ser	Val	Ser	Leu	Trp	Cys	Thr	Ala	Asp	Arg	Ser																																
				565								570			575																																

Lys Tyr Leu Gly Tyr Pro Pro Pro Glu Ile Lys Trp Tyr Lys Asn Gly
355 360 365

Ile Pro Leu Glu Ser Asn His Thr Ile Lys Ala Gly His Val Leu Thr
370 375 380

Ile Met Glu Val Ser Glu Arg Asp Thr Gly Asn Tyr Thr Val Ile Leu
385 390 395 400

Thr Asn Pro Ile Ser Lys Glu Lys Gln Ser His Val Val Ser Leu Val
405 410 415

Val Tyr Val Pro Pro Gln Ile Gly Glu Lys Ser Leu Ile Ser Pro Val
420 425 430

Asp Ser Tyr Gln Tyr Gly Thr Thr Gln Thr Leu Thr Cys Thr Val Tyr
435 440 445

Ala Ile Pro Pro Pro His His Ile His Trp Tyr Trp Gln Leu Glu Glu
450 455 460

Glu Cys Ala Asn Glu Pro Ser Gln Ala Val Ser Val Thr Asn Pro Tyr
465 470 475 480

Pro Cys Glu Glu Trp Arg Ser Val Glu Asp Phe Gln Gly Gly Asn Lys
485 490 495

Ile Glu Val Asn Lys Asn Gln Phe Ala Leu Ile Glu Gly Lys Asn Lys
500 505 510

Thr Val Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr
515 520 525

Lys Cys Glu Ala Val Asn Lys Val Gly Arg Gly Glu Arg Val Ile Ser
530 535 540

Phe His Val Thr Arg Gly Pro Glu Ile Thr Leu Gln Pro Asp Met Gln
545 550 555 560

Pro Thr Glu Gln Glu Ser Val Ser Leu Trp Cys Thr Ala Asp Arg Ser
565 570 575

Thr Phe Glu Asn Leu Thr Trp Tyr Lys Leu Gly Pro Gln Pro Leu Pro
580 585 590

Ile His Val Gly Glu Leu Pro Thr Pro Val Cys Lys Asn Leu Asp Thr
595 600 605

Leu Trp Lys Leu Asn Ala Thr Met Phe Ser Asn Ser Thr Asn Asp Ile
610 615 620

Leu Ile Met Glu Leu Lys Asn Ala Ser Leu Gln Asp Gln Gly Asp Tyr
625 630 635 640

Val Cys Leu Ala Gln Asp Arg Lys Thr Lys Lys Arg His Cys Val Val
645 650 655

Arg Gln Leu Thr Val Leu Glu Arg Val Ala Pro Thr Ile Thr Gly Asn
660 665 670

Leu Glu Asn Gln Thr Thr Ser Ile Gly Glu Ser Ile Glu Val Ser Cys
675 680 685

Thr Ala Ser Gly Asn Pro Pro Pro Gln Ile Met Trp Phe Lys Asp Asn
690 695 700

Glu Thr Leu Val Glu Asp Ser Gly Ile Val Leu Lys Asp Gly Asn Arg
705 710 715 720

Asn Leu Thr Ile Arg Arg Val Arg Lys Glu Asp Glu Gly Leu Tyr Thr
725 730 735

Cys Gln Ala Cys Ser Val Leu Gly Cys Ala Lys Val Glu Ala Phe Phe
740 745 750

Ile Ile Glu Gly Ala Gln Glu Lys Thr Asn Leu Glu Ile Ile Ile Leu
755 760 765

Val Gly Thr Ala Val Ile Ala Met Phe Phe Trp Leu Leu Leu Val Ile
770 775 780

Ile Leu Arg Thr Val Lys Arg Ala Asn Gly Gly Glu Leu Lys Thr Gly
785 790 795 800

Tyr Leu Ser Ile Val Met Asp Pro Asp Glu Leu Pro Leu Asp Glu His
805 810 815

Cys Glu Arg Leu Pro Tyr Asp Ala Ser Lys Trp Glu Phe Pro Arg Asp
820 825 830

Arg Leu Lys Leu Gly Lys Pro Leu Gly Arg Gly Ala Phe Gly Gln Val
835 840 845

Ile Glu Ala Asp Ala Phe Gly Ile Asp Lys Thr Ala Thr Cys Arg Thr
850 855 860

Val Ala Val Lys Met Leu Lys Glu Gly Ala Thr His Ser Glu His Arg
865 870 875 880

Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly His His Leu
885 890 895

Asn Val Val Asn Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu
900 905 910

Met Val Ile Val Glu Phe Cys Lys Phe Gly Asn Leu Ser Thr Tyr Leu
915 920 925

Arg Ser Lys Arg Asn Glu Phe Val Pro Tyr Lys Thr Lys Gly Ala Arg
930 935 940

Phe Arg Gln Gly Lys Asp Tyr Val Gly Ala Ile Pro Val Asp Leu Lys
945 950 955 960

Arg Arg Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly
965 970 975

Phe Val Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Pro
980 985 990

Glu Asp Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr
995 1000 1005

Ser Phe Gln Val Ala Lys Gly Met Glu Phe Leu Ala Ser Arg Lys
1010 1015 1020

Cys	Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Leu	Ser	Glu
1025						1030					1035			
Lys	Asn	Val	Val	Lys	Ile	Cys	Asp	Phe	Gly	Leu	Ala	Arg	Asp	Ile
1040						1045					1050			
Tyr	Lys	Asp	Pro	Asp	Tyr	Val	Arg	Lys	Gly	Asp	Ala	Arg	Leu	Pro
1055						1060					1065			
Leu	Lys	Trp	Met	Ala	Pro	Glu	Thr	Ile	Phe	Asp	Arg	Val	Tyr	Thr
1070						1075					1080			
Ile	Gln	Ser	Asp	Val	Trp	Ser	Phe	Gly	Val	Leu	Leu	Trp	Glu	Ile
1085						1090					1095			
Phe	Ser	Leu	Gly	Ala	Ser	Pro	Tyr	Pro	Gly	Val	Lys	Ile	Asp	Glu
1100						1105					1110			
Glu	Phe	Cys	Arg	Arg	Leu	Lys	Glu	Gly	Thr	Arg	Met	Arg	Ala	Pro
1115						1120					1125			
Asp	Tyr	Thr	Thr	Pro	Glu	Met	Tyr	Gln	Thr	Met	Leu	Asp	Cys	Trp
1130						1135					1140			
His	Gly	Glu	Pro	Ser	Gln	Arg	Pro	Thr	Phe	Ser	Glu	Leu	Val	Glu
1145						1150					1155			
His	Leu	Gly	Asn	Leu	Leu	Gln	Ala	Asn	Ala	Gln	Gln	Asp	Gly	Lys
1160						1165					1170			
Asp	Tyr	Ile	Val	Leu	Pro	Ile	Ser	Glu	Thr	Leu	Ser	Met	Glu	Glu
1175						1180					1185			
Asp	Ser	Gly	Leu	Ser	Leu	Pro	Thr	Ser	Pro	Val	Ser	Cys	Met	Glu
1190						1195					1200			
Glu	Glu	Glu	Val	Cys	Asp	Pro	Lys	Phe	His	Tyr	Asp	Asn	Thr	Ala
1205						1210					1215			
Gly	Ile	Ser	Gln	Tyr	Leu	Gln	Asn	Ser	Lys	Arg	Lys	Ser	Arg	Pro
1220						1225					1230			
Val	Ser	Val	Lys	Thr	Phe	Glu	Asp	Ile	Pro	Leu	Glu	Glu	Pro	Glu

1235		1240		1245
Val Lys	Val Ile Pro Asp Asp	Asn Gln Thr Asp	Ser Gly Met Val	
1250		1255	1260	
Leu Ala	Ser Glu Glu Leu Lys	Thr Leu Glu Asp	Arg Thr Lys Leu	
1265		1270	1275	
Ser Pro	Ser Phe Gly Gly Met	Val Pro Ser Lys	Ser Arg Glu Ser	
1280		1285	1290	
Val Ala	Ser Glu Gly Ser Asn	Gln Thr Ser Gly Tyr	Gln Ser Gly	
1295		1300	1305	
Tyr His	Ser Asp Asp Thr Asp	Thr Thr Val Tyr	Ser Ser Glu Glu	
1310		1315	1320	
Ala Glu	Leu Leu Lys Leu Ile	Glu Ile Gly Val	Gln Thr Gly Ser	
1325		1330	1335	
Thr Ala	Gln Ile Leu Gln Pro	Asp Ser Gly Thr Thr	Leu Ser Ser	
1340		1345	1350	
Pro Pro	Val			
1355				
<210>	21			
<211>	569			
<212>	PRT			
<213>	Homo sapiens			
<400>	21			
Met Lys Val Leu Leu Arg Leu Ile Cys Phe Ile Ala Leu Leu Ile Ser				
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Ser Leu Glu Ala Asp Lys Cys Lys Glu Arg Glu Glu Lys Ile Ile Leu				
	20	25	30	
Val Ser Ser Ala Asn Glu Ile Asp Val Arg Pro Cys Pro Leu Asn Pro				
	35	40	45	
Asn Glu His Lys Gly Thr Ile Thr Trp Tyr Lys Asp Asp Ser Lys Thr				
50	55	60		

Pro Val Ser Thr Glu Gln Ala Ser Arg Ile His Gln His Lys Glu Lys
65 70 75 80

Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly His Tyr Tyr Cys
85 90 95

Val Val Arg Asn Ser Ser Tyr Cys Leu Arg Ile Lys Ile Ser Ala Lys
100 105 110

Phe Val Glu Asn Glu Pro Asn Leu Cys Tyr Asn Ala Gln Ala Ile Phe
115 120 125

Lys Gln Lys Leu Pro Val Ala Gly Asp Gly Gly Leu Val Cys Pro Tyr
130 135 140

Met Glu Phe Phe Lys Asn Glu Asn Asn Glu Leu Pro Lys Leu Gln Trp
145 150 155 160

Tyr Lys Asp Cys Lys Pro Leu Leu Leu Asp Asn Ile His Phe Ser Gly
165 170 175

Val Lys Asp Arg Leu Ile Val Met Asn Val Ala Glu Lys His Arg Gly
180 185 190

Asn Tyr Thr Cys His Ala Ser Tyr Thr Tyr Leu Gly Lys Gln Tyr Pro
195 200 205

Ile Thr Arg Val Ile Glu Phe Ile Thr Leu Glu Glu Asn Lys Pro Thr
210 215 220

Arg Pro Val Ile Val Ser Pro Ala Asn Glu Thr Met Glu Val Asp Leu
225 230 235 240

Gly Ser Gln Ile Gln Leu Ile Cys Asn Val Thr Gly Gln Leu Ser Asp
245 250 255

Ile Ala Tyr Trp Lys Trp Asn Gly Ser Val Ile Asp Glu Asp Asp Pro
260 265 270

Val Leu Gly Glu Asp Tyr Tyr Ser Val Glu Asn Pro Ala Asn Lys Arg
275 280 285

Arg Ser Thr Leu Ile Thr Val Leu Asn Ile Ser Glu Ile Glu Ser Arg
290 295 300

Phe Tyr Lys His Pro Phe Thr Cys Phe Ala Lys Asn Thr His Gly Ile
305 310 315 320

Asp Ala Ala Tyr Ile Gln Leu Ile Tyr Pro Val Thr Asn Phe Gln Lys
325 330 335

His Met Ile Gly Ile Cys Val Thr Leu Thr Val Ile Ile Val Cys Ser
340 345 350

Val Phe Ile Tyr Lys Ile Phe Lys Ile Asp Ile Val Leu Trp Tyr Arg
355 360 365

Asp Ser Cys Tyr Asp Phe Leu Pro Ile Lys Ala Ser Asp Gly Lys Thr
370 375 380

Tyr Asp Ala Tyr Ile Leu Tyr Pro Lys Thr Val Gly Glu Gly Ser Thr
385 390 395 400

Ser Asp Cys Asp Ile Phe Val Phe Lys Val Leu Pro Glu Val Leu Glu
405 410 415

Lys Gln Cys Gly Tyr Lys Leu Phe Ile Tyr Gly Arg Asp Asp Tyr Val
420 425 430

Gly Glu Asp Ile Val Glu Val Ile Asn Glu Asn Val Lys Lys Ser Arg
435 440 445

Arg Leu Ile Ile Ile Leu Val Arg Glu Thr Ser Gly Phe Ser Trp Leu
450 455 460

Gly Gly Ser Ser Glu Glu Gln Ile Ala Met Tyr Asn Ala Leu Val Gln
465 470 475 480

Asp Gly Ile Lys Val Val Leu Leu Glu Leu Glu Lys Ile Gln Asp Tyr
485 490 495

Glu Lys Met Pro Glu Ser Ile Lys Phe Ile Lys Gln Lys His Gly Ala
500 505 510

Ile Arg Trp Ser Gly Asp Phe Thr Gln Gly Pro Gln Ser Ala Lys Thr

515

520

525

Arg Phe Trp Lys Asn Val Arg Tyr His Met Pro Val Gln Arg Arg Ser
 530 535 540

Pro Ser Ser Lys His Gln Leu Leu Ser Pro Ala Thr Lys Glu Lys Leu
 545 550 555 560

Gln Arg Glu Ala His Val Pro Leu Gly
 565

<210> 22

<211> 570

<212> PRT

<213> Homo sapiens

<400> 22

Met Thr Leu Leu Trp Cys Val Val Ser Leu Tyr Phe Tyr Gly Ile Leu
 1 5 10 15

Gln Ser Asp Ala Ser Glu Arg Cys Asp Asp Trp Gly Leu Asp Thr Met
 20 25 30

Arg Gln Ile Gln Val Phe Glu Asp Glu Pro Ala Arg Ile Lys Cys Pro
 35 40 45

Leu Phe Glu His Phe Leu Lys Phe Asn Tyr Ser Thr Ala His Ser Ala
 50 55 60

Gly Leu Thr Leu Ile Trp Tyr Trp Thr Arg Gln Asp Arg Asp Leu Glu
 65 70 75 80

Glu Pro Ile Asn Phe Arg Leu Pro Glu Asn Arg Ile Ser Lys Glu Lys
 85 90 95

Asp Val Leu Trp Phe Arg Pro Thr Leu Leu Asn Asp Thr Gly Asn Tyr
 100 105 110

Thr Cys Met Leu Arg Asn Thr Thr Tyr Cys Ser Lys Val Ala Phe Pro
 115 120 125

Leu Glu Val Val Gln Lys Asp Ser Cys Phe Asn Ser Pro Met Lys Leu
 130 135 140

Pro Val His Lys Leu Tyr Ile Glu Tyr Gly Ile Gln Arg Ile Thr Cys
 145 150 155 160

Pro Asn Val Asp Gly Tyr Phe Pro Ser Ser Val Lys Pro Thr Ile Thr
 165 170 175

Trp Tyr Met Gly Cys Tyr Lys Ile Gln Asn Phe Asn Asn Val Ile Pro
 180 185 190

Glu Gly Met Asn Leu Ser Phe Leu Ile Ala Leu Ile Ser Asn Asn Gly
 195 200 205

Asn Tyr Thr Cys Val Val Thr Tyr Pro Glu Asn Gly Arg Thr Phe His
 210 215 220

Leu Thr Arg Thr Leu Thr Val Lys Val Val Gly Ser Pro Lys Asn Ala
 225 230 235 240

Val Pro Pro Val Ile His Ser Pro Asn Asp His Val Val Tyr Glu Lys
 245 250 255

Glu Pro Gly Glu Glu Leu Leu Ile Pro Cys Thr Val Tyr Phe Ser Phe
 260 265 270

Leu Met Asp Ser Arg Asn Glu Val Trp Trp Thr Ile Asp Gly Lys Lys
 275 280 285

Pro Asp Asp Ile Thr Ile Asp Val Thr Ile Asn Glu Ser Ile Ser His
 290 295 300

Ser Arg Thr Glu Asp Glu Thr Arg Thr Gln Ile Leu Ser Ile Lys Lys
 305 310 315 320

Val Thr Ser Glu Asp Leu Lys Arg Ser Tyr Val Cys His Ala Arg Ser
 325 330 335

Ala Lys Gly Glu Val Ala Lys Ala Ala Lys Val Lys Gln Lys Val Pro
 340 345 350

Ala Pro Arg Tyr Thr Val Glu Leu Ala Cys Gly Phe Gly Ala Thr Val
 355 360 365

Leu Leu Val Val Ile Leu Ile Val Val Tyr His Val Tyr Trp Leu Glu
370 375 380

Met Val Leu Phe Tyr Arg Ala His Phe Gly Thr Asp Glu Thr Ile Leu
385 390 395 400

Asp Gly Lys Glu Tyr Asp Ile Tyr Val Ser Tyr Ala Arg Asn Ala Glu
405 410 415

Glu Glu Glu Phe Val Leu Leu Thr Leu Arg Gly Val Leu Glu Asn Glu
420 425 430

Phe Gly Tyr Lys Leu Cys Ile Phe Asp Arg Asp Ser Leu Pro Gly Gly
435 440 445

Ile Val Thr Asp Glu Thr Leu Ser Phe Ile Gln Lys Ser Arg Arg Leu
450 455 460

Leu Val Val Leu Ser Pro Asn Tyr Val Leu Gln Gly Thr Gln Ala Leu
465 470 475 480

Leu Glu Leu Lys Ala Gly Leu Glu Asn Met Ala Ser Arg Gly Asn Ile
485 490 495

Asn Val Ile Leu Val Gln Tyr Lys Ala Val Lys Glu Thr Lys Val Lys
500 505 510

Glu Leu Lys Arg Ala Lys Thr Val Leu Thr Val Ile Lys Trp Lys Gly
515 520 525

Glu Lys Ser Lys Tyr Pro Gln Gly Arg Phe Trp Lys Gln Leu Gln Val
530 535 540

Ala Met Pro Val Lys Lys Ser Pro Arg Arg Ser Ser Ser Asp Glu Gln
545 550 555 560

Gly Leu Ser Tyr Ser Ser Leu Lys Asn Val
565 570

<210> 23
<211> 1367
<212> PRT
<213> Homo sapiens

<400> 23

Met Lys Ser Gly Ser Gly Gly Gly Ser Pro Thr Ser Leu Trp Gly Leu
1 5 10 15

Leu Phe Leu Ser Ala Ala Leu Ser Leu Trp Pro Thr Ser Gly Glu Ile
20 25 30

Cys Gly Pro Gly Ile Asp Ile Arg Asn Asp Tyr Gln Gln Leu Lys Arg
35 40 45

Leu Glu Asn Cys Thr Val Ile Glu Gly Tyr Leu His Ile Leu Leu Ile
50 55 60

Ser Lys Ala Glu Asp Tyr Arg Ser Tyr Arg Phe Pro Lys Leu Thr Val
65 70 75 80

Ile Thr Glu Tyr Leu Leu Leu Phe Arg Val Ala Gly Leu Glu Ser Leu
85 90 95

Gly Asp Leu Phe Pro Asn Leu Thr Val Ile Arg Gly Trp Lys Leu Phe
100 105 110

Tyr Asn Tyr Ala Leu Val Ile Phe Glu Met Thr Asn Leu Lys Asp Ile
115 120 125

Gly Leu Tyr Asn Leu Arg Asn Ile Thr Arg Gly Ala Ile Arg Ile Glu
130 135 140

Lys Asn Ala Asp Leu Cys Tyr Leu Ser Thr Val Asp Trp Ser Leu Ile
145 150 155 160

Leu Asp Ala Val Ser Asn Asn Tyr Ile Val Gly Asn Lys Pro Pro Lys
165 170 175

Glu Cys Gly Asp Leu Cys Pro Gly Thr Met Glu Glu Lys Pro Met Cys
180 185 190

Glu Lys Thr Thr Ile Asn Asn Glu Tyr Asn Tyr Arg Cys Trp Thr Thr
195 200 205

Asn Arg Cys Gln Lys Met Cys Pro Ser Thr Cys Gly Lys Arg Ala Cys
210 215 220

Thr Glu Asn Asn Glu Cys Cys His Pro Glu Cys Leu Gly Ser Cys Ser
225 230 235 240

Ala Pro Asp Asn Asp Thr Ala Cys Val Ala Cys Arg His Tyr Tyr Tyr
245 250 255

Ala Gly Val Cys Val Pro Ala Cys Pro Pro Asn Thr Tyr Arg Phe Glu
260 265 270

Gly Trp Arg Cys Val Asp Arg Asp Phe Cys Ala Asn Ile Leu Ser Ala
275 280 285

Glu Ser Ser Asp Ser Glu Gly Phe Val Ile His Asp Gly Glu Cys Met
290 295 300

Gln Glu Cys Pro Ser Gly Phe Ile Arg Asn Gly Ser Gln Ser Met Tyr
305 310 315 320

Cys Ile Pro Cys Glu Gly Pro Cys Pro Lys Val Cys Glu Glu Glu Lys
325 330 335

Lys Thr Lys Thr Ile Asp Ser Val Thr Ser Ala Gln Met Leu Gln Gly
340 345 350

Cys Thr Ile Phe Lys Gly Asn Leu Leu Ile Asn Ile Arg Arg Gly Asn
355 360 365

Asn Ile Ala Ser Glu Leu Glu Asn Phe Met Gly Leu Ile Glu Val Val
370 375 380

Thr Gly Tyr Val Lys Ile Arg His Ser His Ala Leu Val Ser Leu Ser
385 390 395 400

Phe Leu Lys Asn Leu Arg Leu Ile Leu Gly Glu Glu Gln Leu Glu Gly
405 410 415

Asn Tyr Ser Phe Tyr Val Leu Asp Asn Gln Asn Leu Gln Gln Leu Trp
420 425 430

Asp Trp Asp His Arg Asn Leu Thr Ile Lys Ala Gly Lys Met Tyr Phe
435 440 445

Ala Phe Asn Pro Lys Leu Cys Val Ser Glu Ile Tyr Arg Met Glu Glu
450 455 460

Val Thr Gly Thr Lys Gly Arg Gln Ser Lys Gly Asp Ile Asn Thr Arg
465 470 475 480

Asn Asn Gly Glu Arg Ala Ser Cys Glu Ser Asp Val Leu His Phe Thr
485 490 495

Ser Thr Thr Thr Ser Lys Asn Arg Ile Ile Ile Thr Trp His Arg Tyr
500 505 510

Arg Pro Pro Asp Tyr Arg Asp Leu Ile Ser Phe Thr Val Tyr Tyr Lys
515 520 525

Glu Ala Pro Phe Lys Asn Val Thr Glu Tyr Asp Gly Gln Asp Ala Cys
530 535 540

Gly Ser Asn Ser Trp Asn Met Val Asp Val Asp Leu Pro Pro Asn Lys
545 550 555 560

Asp Val Glu Pro Gly Ile Leu Leu His Gly Leu Lys Pro Trp Thr Gln
565 570 575

Tyr Ala Val Tyr Val Lys Ala Val Thr Leu Thr Met Val Glu Asn Asp
580 585 590

His Ile Arg Gly Ala Lys Ser Glu Ile Leu Tyr Ile Arg Thr Asn Ala
595 600 605

Ser Val Pro Ser Ile Pro Leu Asp Val Leu Ser Ala Ser Asn Ser Ser
610 615 620

Ser Gln Leu Ile Val Lys Trp Asn Pro Pro Ser Leu Pro Asn Gly Asn
625 630 635 640

Leu Ser Tyr Tyr Ile Val Arg Trp Gln Arg Gln Pro Gln Asp Gly Tyr
645 650 655

Leu Tyr Arg His Asn Tyr Cys Ser Lys Asp Lys Ile Pro Ile Arg Lys
660 665 670

Tyr Ala Asp Gly Thr Ile Asp Ile Glu Glu Val Thr Glu Asn Pro Lys

675

680

685

Thr Glu Val Cys Gly Gly Glu Lys Gly Pro Cys Cys Ala Cys Pro Lys
690 695 700

Thr Glu Ala Glu Lys Gln Ala Glu Lys Glu Glu Ala Glu Tyr Arg Lys
705 710 715 720

Val Phe Glu Asn Phe Leu His Asn Ser Ile Phe Val Pro Arg Pro Glu
725 730 735

Arg Lys Arg Arg Asp Val Met Gln Val Ala Asn Thr Thr Met Ser Ser
740 745 750

Arg Ser Arg Asn Thr Thr Ala Ala Asp Thr Tyr Asn Ile Thr Asp Pro
755 760 765

Glu Glu Leu Glu Thr Glu Tyr Pro Phe Phe Glu Ser Arg Val Asp Asn
770 775 780

Lys Glu Arg Thr Val Ile Ser Asn Leu Arg Pro Phe Thr Leu Tyr Arg
785 790 795 800

Ile Asp Ile His Ser Cys Asn His Glu Ala Glu Lys Leu Gly Cys Ser
805 810 815

Ala Ser Asn Phe Val Phe Ala Arg Thr Met Pro Ala Glu Gly Ala Asp
820 825 830

Asp Ile Pro Gly Pro Val Thr Trp Glu Pro Arg Pro Glu Asn Ser Ile
835 840 845

Phe Leu Lys Trp Pro Glu Pro Glu Asn Pro Asn Gly Leu Ile Leu Met
850 855 860

Tyr Glu Ile Lys Tyr Gly Ser Gln Val Glu Asp Gln Arg Glu Cys Val
865 870 875 880

Ser Arg Gln Glu Tyr Arg Lys Tyr Gly Gly Ala Lys Leu Asn Arg Leu
885 890 895

Asn Pro Gly Asn Tyr Thr Ala Arg Ile Gln Ala Thr Ser Leu Ser Gly
900 905 910

Asn Gly Ser Trp Thr Asp Pro Val Phe Phe Tyr Val Gln Ala Lys Thr
 915 920 925

Gly Tyr Glu Asn Phe Ile His Leu Ile Ile Ala Leu Pro Val Ala Val
 930 935 940

Leu Leu Ile Val Gly Gly Leu Val Ile Met Leu Tyr Val Phe His Arg
 945 950 955 960

Lys Arg Asn Asn Ser Arg Leu Gly Asn Gly Val Leu Tyr Ala Ser Val
 965 970 975

Asn Pro Glu Tyr Phe Ser Ala Ala Asp Val Tyr Val Pro Asp Glu Trp
 980 985 990

Glu Val Ala Arg Glu Lys Ile Thr Met Ser Arg Glu Leu Gly Gln Gly
 995 1000 1005

Ser Phe Gly Met Val Tyr Glu Gly Val Ala Lys Gly Val Val Lys
 1010 1015 1020

Asp Glu Pro Glu Thr Arg Val Ala Ile Lys Thr Val Asn Glu Ala
 1025 1030 1035

Ala Ser Met Arg Glu Arg Ile Glu Phe Leu Asn Glu Ala Ser Val
 1040 1045 1050

Met Lys Glu Phe Asn Cys His His Val Val Arg Leu Leu Gly Val
 1055 1060 1065

Val Ser Gln Gly Gln Pro Thr Leu Val Ile Met Glu Leu Met Thr
 1070 1075 1080

Arg Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Met
 1085 1090 1095

Glu Asn Asn Pro Val Leu Ala Pro Pro Ser Leu Ser Lys Met Ile
 1100 1105 1110

Gln Met Ala Gly Glu Ile Ala Asp Gly Met Ala Tyr Leu Asn Ala
 1115 1120 1125

Asn Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met Val		
1130	1135	1140
Ala Glu Asp Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg		
1145	1150	1155
Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Gly Lys Gly Leu		
1160	1165	1170
Leu Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Val		
1175	1180	1185
Phe Thr Thr Tyr Ser Asp Val Trp Ser Phe Gly Val Val Leu Trp		
1190	1195	1200
Glu Ile Ala Thr Leu Ala Glu Gln Pro Tyr Gln Gly Leu Ser Asn		
1205	1210	1215
Glu Gln Val Leu Arg Phe Val Met Glu Gly Gly Leu Leu Asp Lys		
1220	1225	1230
Pro Asp Asn Cys Pro Asp Met Leu Phe Glu Leu Met Arg Met Cys		
1235	1240	1245
Trp Gln Tyr Asn Pro Lys Met Arg Pro Ser Phe Leu Glu Ile Ile		
1250	1255	1260
Ser Ser Ile Lys Glu Glu Met Glu Pro Gly Phe Arg Glu Val Ser		
1265	1270	1275
Phe Tyr Tyr Ser Glu Glu Asn Lys Leu Pro Glu Pro Glu Glu Leu		
1280	1285	1290
Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp Pro Ser		
1295	1300	1305
Ala Ser Ser Ser Ser Leu Pro Leu Pro Asp Arg His Ser Gly His		
1310	1315	1320
Lys Ala Glu Asn Gly Pro Gly Pro Gly Val Leu Val Leu Arg Ala		
1325	1330	1335

Ser Phe Asp Glu Arg Gln Pro Tyr Ala His Met Asn Gly Gly Arg
 1340 1345 1350

Lys Asn Glu Arg Ala Leu Pro Leu Pro Gln Ser Ser Thr Cys
 1355 1360 1365

<210> 24
 <211> 825
 <212> PRT
 <213> Homo sapiens

<400> 24

Met Gly Trp Leu Cys Ser Gly Leu Leu Phe Pro Val Ser Cys Leu Val
 1 5 10 15

Leu Leu Gln Val Ala Ser Ser Gly Asn Met Lys Val Leu Gln Glu Pro
 20 25 30

Thr Cys Val Ser Asp Tyr Met Ser Ile Ser Thr Cys Glu Trp Lys Met
 35 40 45

Asn Gly Pro Thr Asn Cys Ser Thr Glu Leu Arg Leu Leu Tyr Gln Leu
 50 55 60

Val Phe Leu Leu Ser Glu Ala His Thr Cys Ile Pro Glu Asn Asn Gly
 65 70 75 80

Gly Ala Gly Cys Val Cys His Leu Leu Met Asp Asp Val Val Ser Ala
 85 90 95

Asp Asn Tyr Thr Leu Asp Leu Trp Ala Gly Gln Gln Leu Leu Trp Lys
 100 105 110

Gly Ser Phe Lys Pro Ser Glu His Val Lys Pro Arg Ala Pro Gly Asn
 115 120 125

Leu Thr Val His Thr Asn Val Ser Asp Thr Leu Leu Leu Thr Trp Ser
 130 135 140

Asn Pro Tyr Pro Pro Asp Asn Tyr Leu Tyr Asn His Leu Thr Tyr Ala
 145 150 155 160

Val Asn Ile Trp Ser Glu Asn Asp Pro Ala Asp Phe Arg Ile Tyr Asn
 165 170 175

Val Thr Tyr Leu Glu Pro Ser Leu Arg Ile Ala Ala Ser Thr Leu Lys
180 185 190

Ser Gly Ile Ser Tyr Arg Ala Arg Val Arg Ala Trp Ala Gln Cys Tyr
195 200 205

Asn Thr Thr Trp Ser Glu Trp Ser Pro Ser Thr Lys Trp His Asn Ser
210 215 220

Tyr Arg Glu Pro Phe Glu Gln His Leu Leu Leu Gly Val Ser Val Ser
225 230 235 240

Cys Ile Val Ile Leu Ala Val Cys Leu Leu Cys Tyr Val Ser Ile Thr
245 250 255

Lys Ile Lys Lys Glu Trp Trp Asp Gln Ile Pro Asn Pro Ala Arg Ser
260 265 270

Arg Leu Val Ala Ile Ile Ile Gln Asp Ala Gln Gly Ser Gln Trp Glu
275 280 285

Lys Arg Ser Arg Gly Gln Glu Pro Ala Lys Cys Pro His Trp Lys Asn
290 295 300

Cys Leu Thr Lys Leu Leu Pro Cys Phe Leu Glu His Asn Met Lys Arg
305 310 315 320

Asp Glu Asp Pro His Lys Ala Ala Lys Glu Met Pro Phe Gln Gly Ser
325 330 335

Gly Lys Ser Ala Trp Cys Pro Val Glu Ile Ser Lys Thr Val Leu Trp
340 345 350

Pro Glu Ser Ile Ser Val Val Arg Cys Val Glu Leu Phe Glu Ala Pro
355 360 365

Val Glu Cys Glu Glu Glu Glu Glu Val Glu Glu Glu Lys Gly Ser Phe
370 375 380

Cys Ala Ser Pro Glu Ser Ser Arg Asp Asp Phe Gln Glu Gly Arg Glu
385 390 395 400

Gly Ile Val Ala Arg Leu Thr Glu Ser Leu Phe Leu Asp Leu Leu Gly
405 410 415

Glu Glu Asn Gly Gly Phe Cys Gln Gln Asp Met Gly Glu Ser Cys Leu
420 425 430

Leu Pro Pro Ser Gly Ser Thr Ser Ala His Met Pro Trp Asp Glu Phe
435 440 445

Pro Ser Ala Gly Pro Lys Glu Ala Pro Pro Trp Gly Lys Glu Gln Pro
450 455 460

Leu His Leu Glu Pro Ser Pro Pro Ala Ser Pro Thr Gln Ser Pro Asp
465 470 475 480

Asn Leu Thr Cys Thr Glu Thr Pro Leu Val Ile Ala Gly Asn Pro Ala
485 490 495

Tyr Arg Ser Phe Ser Asn Ser Leu Ser Gln Ser Pro Cys Pro Arg Glu
500 505 510

Leu Gly Pro Asp Pro Leu Leu Ala Arg His Leu Glu Glu Val Glu Pro
515 520 525

Glu Met Pro Cys Val Pro Gln Leu Ser Glu Pro Thr Thr Val Pro Gln
530 535 540

Pro Glu Pro Glu Thr Trp Glu Gln Ile Leu Arg Arg Asn Val Leu Gln
545 550 555 560

His Gly Ala Ala Ala Ala Pro Val Ser Ala Pro Thr Ser Gly Tyr Gln
565 570 575

Glu Phe Val His Ala Val Glu Gln Gly Gly Thr Gln Ala Ser Ala Val
580 585 590

Val Gly Leu Gly Pro Pro Gly Glu Ala Gly Tyr Lys Ala Phe Ser Ser
595 600 605

Leu Leu Ala Ser Ser Ala Val Ser Pro Glu Lys Cys Gly Phe Gly Ala
610 615 620

Ser Ser Gly Glu Glu Gly Tyr Lys Pro Phe Gln Asp Leu Ile Pro Gly
625 630 635 640

Cys Pro Gly Asp Pro Ala Pro Val Pro Val Pro Leu Phe Thr Phe Gly
645 650 655

Leu Asp Arg Glu Pro Pro Arg Ser Pro Gln Ser Ser His Leu Pro Ser
660 665 670

Ser Ser Pro Glu His Leu Gly Leu Glu Pro Gly Glu Lys Val Glu Asp
675 680 685

Met Pro Lys Pro Pro Leu Pro Gln Glu Gln Ala Thr Asp Pro Leu Val
690 695 700

Asp Ser Leu Gly Ser Gly Ile Val Tyr Ser Ala Leu Thr Cys His Leu
705 710 715 720

Cys Gly His Leu Lys Gln Cys His Gly Gln Glu Asp Gly Gly Gln Thr
725 730 735

Pro Val Met Ala Ser Pro Cys Cys Gly Cys Cys Cys Gly Asp Arg Ser
740 745 750

Ser Pro Pro Thr Thr Pro Leu Arg Ala Pro Asp Pro Ser Pro Gly Gly
755 760 765

Val Pro Leu Glu Ala Ser Leu Cys Pro Ala Ser Leu Ala Pro Ser Gly
770 775 780

Ile Ser Glu Lys Ser Lys Ser Ser Ser Ser Phe His Pro Ala Pro Gly
785 790 795 800

Asn Ala Gln Ser Ser Ser Gln Thr Pro Lys Ile Val Asn Phe Val Ser
805 810 815

Val Gly Pro Thr Tyr Met Arg Val Ser
820 825

<210> 25
<211> 569
<212> PRT
<213> Homo sapiens

<400> 25

Met Lys Val Leu Leu Arg Leu Ile Cys Phe Ile Ala Leu Leu Ile Ser
1 5 10 15

Ser Leu Glu Ala Asp Lys Cys Lys Glu Arg Glu Glu Lys Ile Ile Leu
20 25 30

Val Ser Ser Ala Asn Glu Ile Asp Val Arg Pro Cys Pro Leu Asn Pro
35 40 45

Asn Glu His Lys Gly Thr Ile Thr Trp Tyr Lys Asp Asp Ser Lys Thr
50 55 60

Pro Val Ser Thr Glu Gln Ala Ser Arg Ile His Gln His Lys Glu Lys
65 70 75 80

Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly His Tyr Tyr Cys
85 90 95

Val Val Arg Asn Ser Ser Tyr Cys Leu Arg Ile Lys Ile Ser Ala Lys
100 105 110

Phe Val Glu Asn Glu Pro Asn Leu Cys Tyr Asn Ala Gln Ala Ile Phe
115 120 125

Lys Gln Lys Leu Pro Val Ala Gly Asp Gly Gly Leu Val Cys Pro Tyr
130 135 140

Met Glu Phe Phe Lys Asn Glu Asn Asn Glu Leu Pro Lys Leu Gln Trp
145 150 155 160

Tyr Lys Asp Cys Lys Pro Leu Leu Leu Asp Asn Ile His Phe Ser Gly
165 170 175

Val Lys Asp Arg Leu Ile Val Met Asn Val Ala Glu Lys His Arg Gly
180 185 190

Asn Tyr Thr Cys His Ala Ser Tyr Thr Tyr Leu Gly Lys Gln Tyr Pro
195 200 205

Ile Thr Arg Val Ile Glu Phe Ile Thr Leu Glu Glu Asn Lys Pro Thr
210 215 220

Arg Pro Val Ile Val Ser Pro Ala Asn Glu Thr Met Glu Val Asp Leu
225 230 235 240

Gly Ser Gln Ile Gln Leu Ile Cys Asn Val Thr Gly Gln Leu Ser Asp
245 250 255

Ile Ala Tyr Trp Lys Trp Asn Gly Ser Val Ile Asp Glu Asp Asp Pro
260 265 270

Val Leu Gly Glu Asp Tyr Tyr Ser Val Glu Asn Pro Ala Asn Lys Arg
275 280 285

Arg Ser Thr Leu Ile Thr Val Leu Asn Ile Ser Glu Ile Glu Ser Arg
290 295 300

Phe Tyr Lys His Pro Phe Thr Cys Phe Ala Lys Asn Thr His Gly Ile
305 310 315 320

Asp Ala Ala Tyr Ile Gln Leu Ile Tyr Pro Val Thr Asn Phe Gln Lys
325 330 335

His Met Ile Gly Ile Cys Val Thr Leu Thr Val Ile Ile Val Cys Ser
340 345 350

Val Phe Ile Tyr Lys Ile Phe Lys Ile Asp Ile Val Leu Trp Tyr Arg
355 360 365

Asp Ser Cys Tyr Asp Phe Leu Pro Ile Lys Ala Ser Asp Gly Lys Thr
370 375 380

Tyr Asp Ala Tyr Ile Leu Tyr Pro Lys Thr Val Gly Glu Gly Ser Thr
385 390 395 400

Ser Asp Cys Asp Ile Phe Val Phe Lys Val Leu Pro Glu Val Leu Glu
405 410 415

Lys Gln Cys Gly Tyr Lys Leu Phe Ile Tyr Gly Arg Asp Asp Tyr Val
420 425 430

Gly Glu Asp Ile Val Glu Val Ile Asn Glu Asn Val Lys Lys Ser Arg
435 440 445

Arg Leu Ile Ile Ile Leu Val Arg Glu Thr Ser Gly Phe Ser Trp Leu
450 455 460

Gly Gly Ser Ser Glu Glu Gln Ile Ala Met Tyr Asn Ala Leu Val Gln
465 470 475 480

Asp Gly Ile Lys Val Val Leu Leu Glu Leu Glu Lys Ile Gln Asp Tyr
485 490 495

Glu Lys Met Pro Glu Ser Ile Lys Phe Ile Lys Gln Lys His Gly Ala
500 505 510

Ile Arg Trp Ser Gly Asp Phe Thr Gln Gly Pro Gln Ser Ala Lys Thr
515 520 525

Arg Phe Trp Lys Asn Val Arg Tyr His Met Pro Val Gln Arg Arg Ser
530 535 540

Pro Ser Ser Lys His Gln Leu Leu Ser Pro Ala Thr Lys Glu Lys Leu
545 550 555 560

Gln Arg Glu Ala His Val Pro Leu Gly
565

<210> 26
<211> 576
<212> PRT
<213> Mus sp.

<400> 26

Met Glu Asn Met Lys Val Leu Leu Gly Leu Ile Cys Leu Met Val Pro
1 5 10 15

Leu Leu Ser Leu Glu Ile Asp Val Cys Thr Glu Tyr Pro Asn Gln Ile
20 25 30

Val Leu Phe Leu Ser Val Asn Glu Ile Asp Ile Arg Lys Cys Pro Leu
35 40 45

Thr Pro Asn Lys Met His Gly Asp Thr Ile Ile Trp Tyr Lys Asn Asp
50 55 60

Ser Lys Thr Pro Ile Ser Ala Asp Arg Asp Ser Arg Ile His Gln Gln
65 70 75 80

Asn Glu His Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly Tyr
85 90 95

Tyr Tyr Cys Ile Val Arg Asn Ser Thr Tyr Cys Leu Lys Thr Lys Val
100 105 110

Thr Val Thr Val Leu Glu Asn Asp Pro Gly Leu Cys Tyr Ser Thr Gln
115 120 125

Ala Thr Phe Pro Gln Arg Leu His Ile Ala Gly Asp Gly Ser Leu Val
130 135 140

Cys Pro Tyr Val Ser Tyr Phe Lys Asp Glu Asn Asn Glu Leu Pro Glu
145 150 155 160

Val Gln Trp Tyr Lys Asn Cys Lys Pro Leu Leu Leu Asp Asn Val Ser
165 170 175

Phe Phe Gly Val Lys Asp Lys Leu Leu Val Arg Asn Val Ala Glu Glu
180 185 190

His Arg Gly Asp Tyr Ile Cys Arg Met Ser Tyr Thr Phe Arg Gly Lys
195 200 205

Gln Tyr Pro Val Thr Arg Val Ile Gln Phe Ile Thr Ile Asp Glu Asn
210 215 220

Lys Arg Asp Arg Pro Val Ile Leu Ser Pro Arg Asn Glu Thr Ile Glu
225 230 235 240

Ala Asp Pro Gly Ser Met Ile Gln Leu Ile Cys Asn Val Thr Gly Gln
245 250 255

Phe Ser Asp Leu Val Tyr Trp Lys Trp Asn Gly Ser Glu Ile Glu Trp
260 265 270

Asn Asp Pro Phe Leu Ala Glu Asp Tyr Gln Phe Val Glu His Pro Ser
275 280 285

Thr Lys Arg Lys Tyr Thr Leu Ile Thr Thr Leu Asn Ile Ser Glu Val
290 295 300

Lys Ser Gln Phe Tyr Arg Tyr Pro Phe Ile Cys Val Val Lys Asn Thr
305 310 315 320

Asn Ile Phe Glu Ser Ala His Val Gln Leu Ile Tyr Pro Val Pro Asp
325 330 335

Phe Lys Asn Tyr Leu Ile Gly Gly Phe Ile Ile Leu Thr Ala Thr Ile
340 345 350

Val Cys Cys Val Cys Ile Tyr Lys Val Phe Lys Val Asp Ile Val Leu
355 360 365

Trp Tyr Arg Asp Ser Cys Ser Gly Phe Leu Pro Ser Lys Ala Ser Asp
370 375 380

Gly Lys Thr Tyr Asp Ala Tyr Ile Leu Tyr Pro Lys Thr Leu Gly Glu
385 390 395 400

Gly Ser Phe Ser Asp Leu Asp Thr Phe Val Phe Lys Leu Leu Pro Glu
405 410 415

Val Leu Glu Gly Gln Phe Gly Tyr Lys Leu Phe Ile Tyr Gly Arg Asp
420 425 430

Asp Tyr Val Gly Glu Asp Thr Ile Glu Val Thr Asn Glu Asn Val Lys
435 440 445

Lys Ser Arg Arg Leu Ile Ile Ile Leu Val Arg Asp Met Gly Gly Phe
450 455 460

Ser Trp Leu Gly Gln Ser Ser Glu Glu Gln Ile Ala Ile Tyr Asn Ala
465 470 475 480

Leu Ile Gln Glu Gly Ile Lys Ile Val Leu Leu Glu Leu Glu Lys Ile
485 490 495

Gln Asp Tyr Glu Lys Met Pro Asp Ser Ile Gln Phe Ile Lys Gln Lys
500 505 510

His Gly Val Ile Cys Trp Ser Gly Asp Phe Gln Glu Arg Pro Gln Ser
515 520 525

Ala Lys Thr Arg Phe Trp Lys Asn Leu Arg Tyr Gln Met Pro Ala Gln
530 535 540

Arg Arg Ser Pro Leu Ser Lys His Arg Leu Leu Thr Leu Asp Pro Val
545 550 555 560

Arg Asp Thr Lys Glu Lys Leu Pro Ala Ala Thr His Leu Pro Leu Gly
565 570 575

<210> 27
<211> 576
<212> PRT
<213> Rattus sp.

<400> 27

Met Glu Asn Met Lys Val Leu Leu Gly Phe Ile Cys Leu Ile Val Pro
1 5 10 15

Leu Leu Ser Leu Glu Thr Asp Lys Cys Thr Glu Tyr Pro Asn Glu Val
20 25 30

Ile Ser Phe Ser Ser Val Asn Glu Ile Asp Ile Arg Ser Cys Pro Leu
35 40 45

Thr Pro Asn Glu Met His Gly Gly Thr Ile Ile Trp Tyr Lys Asn Asp
50 55 60

Ser Lys Thr Pro Ile Ser Ala Asp Lys Asp Ser Arg Ile His Gln Gln
65 70 75 80

Asn Glu His Leu Trp Phe Val Pro Ala Lys Met Glu Asp Ser Gly Tyr
85 90 95

Tyr Tyr Cys Ile Met Arg Asn Ser Thr Tyr Cys Leu Lys Thr Lys Ile
100 105 110

Thr Met Ser Val Leu Glu Asn Asp Pro Gly Leu Cys Tyr Asn Thr Gln
115 120 125

Ala Ser Phe Ile Gln Arg Leu His Val Ala Gly Asp Gly Ser Leu Val
130 135 140

Cys Pro Tyr Leu Asp Phe Phe Lys Asp Glu Asn Asn Glu Leu Pro Lys
145 150 155 160

Val Gln Trp Tyr Lys Asn Cys Lys Pro Leu Pro Leu Asp Asp Gly Asn
165 170 175

Phe Phe Gly Phe Lys Asn Lys Leu Met Val Met Asn Val Ala Glu Glu
180 185 190

His Arg Gly Asn Tyr Thr Cys Arg Thr Ser Tyr Thr Tyr Gln Gly Lys
195 200 205

Gln Tyr Pro Val Thr Arg Val Ile Thr Phe Ile Thr Ile Asp Asp Ser
210 215 220

Lys Arg Asp Arg Pro Val Ile Met Ser Pro Arg Asn Glu Thr Met Glu
225 230 235 240

Ala Asp Pro Gly Ser Thr Ile Gln Leu Ile Cys Asn Val Thr Gly Gln
245 250 255

Phe Thr Asp Leu Val Tyr Trp Lys Trp Asn Gly Ser Glu Ile Glu Trp
260 265 270

Asp Asp Pro Ile Leu Ala Glu Asp Tyr Gln Phe Leu Glu His Pro Ser
275 280 285

Ala Lys Arg Lys Tyr Thr Leu Ile Thr Thr Leu Asn Val Ser Glu Val
290 295 300

Lys Ser Gln Phe Tyr Arg Tyr Pro Phe Ile Cys Phe Val Lys Asn Thr
305 310 315 320

His Ile Leu Glu Thr Ala His Val Arg Leu Val Tyr Pro Val Pro Asp
325 330 335

Phe Lys Asn Tyr Leu Ile Gly Gly Phe Ala Ile Phe Thr Ala Thr Ala
340 345 350

Val Phe Cys Ala Cys Ile Tyr Lys Val Phe Lys Val Asp Ile Val Leu
355 360 365

Trp Tyr Arg Asp Ser Cys Ser Asp Phe Leu Pro Arg Lys Ala Ser Asp
370 375 380

Gly Arg Thr Tyr Asp Ala Tyr Val Leu Tyr Pro Lys Thr Tyr Gly Glu
385 390 395 400

Gly Ser Phe Ala Tyr Leu Asp Thr Phe Val Phe Lys Leu Leu Pro Glu
405 410 415

Val Leu Glu Gly Gln Phe Gly Tyr Lys Leu Phe Ile Cys Gly Arg Asp
420 425 430

Asp Tyr Val Gly Glu Asp Thr Ile Glu Val Thr Asn Glu Asn Val Lys
435 440 445

Arg Ser Arg Arg Leu Ile Ile Ile Leu Val Arg Asp Met Gly Ser Phe
450 455 460

Ser Cys Leu Gly Gln Ser Ser Glu Glu Gln Ile Ala Ile Tyr Asp Ala
465 470 475 480

Leu Ile Arg Glu Gly Ile Lys Ile Ile Leu Leu Glu Leu Glu Lys Ile
485 490 495

Gln Asp Tyr Glu Lys Met Pro Glu Ser Ile Gln Phe Ile Lys Gln Lys
500 505 510

His Gly Ala Ile Cys Trp Ser Gly Asp Phe Lys Glu Arg Pro Gln Ser
515 520 525

Ala Lys Thr Arg Phe Trp Lys Asn Leu Arg Tyr Gln Met Pro Ala Gln
530 535 540

Arg Arg Ser Pro Leu Ser Lys His His Leu Leu Thr Leu Asp Pro Val
545 550 555 560

Leu Asp Thr Lys Glu Lys Leu Gln Ala Glu Thr His Leu Pro Leu Gly
565 570 575

<210> 28
<211> 555
<212> PRT
<213> Equus sp.

<400> 28

Met His Lys Met Thr Ser Thr Phe Leu Leu Ile Gly His Leu Ile Leu

1	5	10	15
Leu Ile Pro	Leu Phe Ser	Ala Glu Glu Cys Val	Ile Cys Asn Tyr Phe
	20	25	30
Val Leu Val	Gly Glu Pro Thr	Ala Ile Ser Cys Pro	Val Ile Thr Leu
	35	40	45
Pro Met Leu	His Ser Asp Tyr	Asn Leu Thr Trp Tyr	Arg Asn Gly Ser
	50	55	60
Asn Met Pro	Ile Thr Thr Glu	Arg Arg Ala Arg	Ile His Gln Arg Lys
	65	70	75
Gly Leu Leu	Trp Phe Ile Pro	Ala Ala Leu Glu	Asp Ser Gly Leu Tyr
	85	90	95
Glu Cys Glu	Val Arg Ser Leu	Asn Arg Ser Lys	Gln Lys Ile Ile Asn
	100	105	110
Leu Lys Val	Phe Lys Asn Asp	Asn Gly Leu Cys	Phe Asn Gly Glu Met
	115	120	125
Lys Tyr Asp	Gln Ile Val Lys	Ser Ala Asn Ala	Gly Lys Ile Ile Cys
	130	135	140
Pro Asp Leu	Glu Asn Phe Lys	Asp Glu Asp Asn	Ile Asn Pro Glu Ile
	145	150	155
His Trp Tyr	Lys Glu Cys Lys	Ser Gly Phe Leu	Glu Asp Lys Arg Leu
	165	170	175
Val Leu Ala	Glu Gly Glu Asn	Ala Ile Leu Ile	Leu Asn Val Thr Ile
	180	185	190
Gln Asp Lys	Gly Asn Tyr Thr	Cys Arg Met Val	Tyr Thr Tyr Met Gly
	195	200	205
Lys Gln Tyr	Asn Val Ser Arg	Thr Met Asn Leu	Glu Val Lys Glu Ser
	210	215	220
Pro Leu Lys	Met Arg Pro Glu	Phe Ile Tyr Pro	Asn Asn Asn Thr Ile
	225	230	235
			240

Glu Val Glu Leu Gly Ser His Val Val Met Glu Cys Asn Val Ser Ser
245 250 255

Gly Val Tyr Gly Leu Leu Pro Tyr Trp Gln Val Asn Asp Glu Asp Val
260 265 270

Asp Ser Phe Asp Ser Thr Tyr Arg Glu Gln Phe Tyr Glu Glu Gly Met
275 280 285

Pro His Gly Ile Ala Val Ser Gly Thr Lys Phe Asn Ile Ser Glu Val
290 295 300

Lys Leu Lys Asp Tyr Ala Tyr Lys Phe Phe Cys His Phe Ile Tyr Asp
305 310 315 320

Ser Gln Glu Phe Thr Ser Tyr Ile Lys Leu Glu His Pro Val Gln Asn
325 330 335

Ile Arg Gly Tyr Leu Ile Gly Gly Gly Ile Ser Leu Ile Phe Leu Leu
340 345 350

Phe Leu Ile Leu Ile Val Tyr Lys Ile Phe Lys Ile Asp Ile Val Leu
355 360 365

Trp Tyr Arg Ser Ser Cys His Pro Leu Leu Gly Lys Lys Val Ser Asp
370 375 380

Gly Lys Ile Tyr Asp Ala Tyr Val Leu Tyr Pro Lys Asn Arg Glu Ser
385 390 395 400

Cys Leu Tyr Ser Ser Asp Ile Phe Ala Leu Lys Ile Leu Pro Glu Val
405 410 415

Leu Glu Arg Gln Cys Gly Tyr Asn Leu Phe Ile Phe Gly Arg Asn Asp
420 425 430

Leu Ala Gly Glu Ala Val Ile Asp Val Thr Asp Glu Lys Ile His Gln
435 440 445

Ser Arg Arg Val Ile Ile Ile Leu Val Pro Glu Pro Ser Cys Tyr Gly
450 455 460

Ile Leu Glu Asp Ala Ser Glu Lys His Leu Ala Val Tyr Asn Ala Leu
465 470 475 480

Ile Gln Asp Gly Ile Lys Ile Ile Leu Ile Glu Leu Glu Lys Ile Glu
485 490 495

Asp Tyr Ala Asn Met Pro Glu Ser Ile Lys Tyr Val Lys Gln Lys Tyr
500 505 510

Gly Ala Ile Arg Trp Thr Gly Asp Phe Ser Glu Arg Ser His Ser Ala
515 520 525

Ser Thr Arg Phe Trp Lys Lys Val Arg Tyr His Met Pro Ser Arg Lys
530 535 540

His Gly Ser Ser Ser Gly Phe His Leu Ser Ser
545 550 555

<210> 29
<211> 825
<212> PRT
<213> Homo sapiens

<400> 29

Met Gly Trp Leu Cys Ser Gly Leu Leu Phe Pro Val Ser Cys Leu Val
1 5 10 15

Leu Leu Gln Val Ala Ser Ser Gly Asn Met Lys Val Leu Gln Glu Pro
20 25 30

Thr Cys Val Ser Asp Tyr Met Ser Ile Ser Thr Cys Glu Trp Lys Met
35 40 45

Asn Gly Pro Thr Asn Cys Ser Thr Glu Leu Arg Leu Leu Tyr Gln Leu
50 55 60

Val Phe Leu Leu Ser Glu Ala His Thr Cys Ile Pro Glu Asn Asn Gly
65 70 75 80

Gly Ala Gly Cys Val Cys His Leu Leu Met Asp Asp Val Val Ser Ala
85 90 95

Asp Asn Tyr Thr Leu Asp Leu Trp Ala Gly Gln Gln Leu Leu Trp Lys

100	105	110
Gly Ser Phe Lys Pro Ser Glu His Val Lys Pro Arg Ala Pro Gly Asn 115 120 125		
Leu Thr Val His Thr Asn Val Ser Asp Thr Leu Leu Leu Thr Trp Ser 130 135 140		
Asn Pro Tyr Pro Pro Asp Asn Tyr Leu Tyr Asn His Leu Thr Tyr Ala 145 150 155 160		
Val Asn Ile Trp Ser Glu Asn Asp Pro Ala Asp Phe Arg Ile Tyr Asn 165 170 175		
Val Thr Tyr Leu Glu Pro Ser Leu Arg Ile Ala Ala Ser Thr Leu Lys 180 185 190		
Ser Gly Ile Ser Tyr Arg Ala Arg Val Arg Ala Trp Ala Gln Cys Tyr 195 200 205		
Asn Thr Thr Trp Ser Glu Trp Ser Pro Ser Thr Lys Trp His Asn Ser 210 215 220		
Tyr Arg Glu Pro Phe Glu Gln His Leu Leu Leu Gly Val Ser Val Ser 225 230 235 240		
Cys Ile Val Ile Leu Ala Val Cys Leu Leu Cys Tyr Val Ser Ile Thr 245 250 255		
Lys Ile Lys Lys Glu Trp Trp Asp Gln Ile Pro Asn Pro Ala Arg Ser 260 265 270		
Arg Leu Val Ala Ile Ile Ile Gln Asp Ala Gln Gly Ser Gln Trp Glu 275 280 285		
Lys Arg Ser Arg Gly Gln Glu Pro Ala Lys Cys Pro His Trp Lys Asn 290 295 300		
Cys Leu Thr Lys Leu Leu Pro Cys Phe Leu Glu His Asn Met Lys Arg 305 310 315 320		
Asp Glu Asp Pro His Lys Ala Ala Lys Glu Met Pro Phe Gln Gly Ser 325 330 335		

Gly Lys Ser Ala Trp Cys Pro Val Glu Ile Ser Lys Thr Val Leu Trp
340 345 350

Pro Glu Ser Ile Ser Val Val Arg Cys Val Glu Leu Phe Glu Ala Pro
355 360 365

Val Glu Cys Glu Glu Glu Glu Glu Val Glu Glu Glu Lys Gly Ser Phe
370 375 380

Cys Ala Ser Pro Glu Ser Ser Arg Asp Asp Phe Gln Glu Gly Arg Glu
385 390 395 400

Gly Ile Val Ala Arg Leu Thr Glu Ser Leu Phe Leu Asp Leu Leu Gly
405 410 415

Glu Glu Asn Gly Gly Phe Cys Gln Gln Asp Met Gly Glu Ser Cys Leu
420 425 430

Leu Pro Pro Ser Gly Ser Thr Ser Ala His Met Pro Trp Asp Glu Phe
435 440 445

Pro Ser Ala Gly Pro Lys Glu Ala Pro Pro Trp Gly Lys Glu Gln Pro
450 455 460

Leu His Leu Glu Pro Ser Pro Pro Ala Ser Pro Thr Gln Ser Pro Asp
465 470 475 480

Asn Leu Thr Cys Thr Glu Thr Pro Leu Val Ile Ala Gly Asn Pro Ala
485 490 495

Tyr Arg Ser Phe Ser Asn Ser Leu Ser Gln Ser Pro Cys Pro Arg Glu
500 505 510

Leu Gly Pro Asp Pro Leu Leu Ala Arg His Leu Glu Glu Val Glu Pro
515 520 525

Glu Met Pro Cys Val Pro Gln Leu Ser Glu Pro Thr Thr Val Pro Gln
530 535 540

Pro Glu Pro Glu Thr Trp Glu Gln Ile Leu Arg Arg Asn Val Leu Gln
545 550 555 560

His Gly Ala Ala Ala Ala Pro Val Ser Ala Pro Thr Ser Gly Tyr Gln
565 570 575

Glu Phe Val His Ala Val Glu Gln Gly Gly Thr Gln Ala Ser Ala Val
580 585 590

Val Gly Leu Gly Pro Pro Gly Glu Ala Gly Tyr Lys Ala Phe Ser Ser
595 600 605

Leu Leu Ala Ser Ser Ala Val Ser Pro Glu Lys Cys Gly Phe Gly Ala
610 615 620

Ser Ser Gly Glu Glu Gly Tyr Lys Pro Phe Gln Asp Leu Ile Pro Gly
625 630 635 640

Cys Pro Gly Asp Pro Ala Pro Val Pro Val Pro Leu Phe Thr Phe Gly
645 650 655

Leu Asp Arg Glu Pro Pro Arg Ser Pro Gln Ser Ser His Leu Pro Ser
660 665 670

Ser Ser Pro Glu His Leu Gly Leu Glu Pro Gly Glu Lys Val Glu Asp
675 680 685

Met Pro Lys Pro Pro Leu Pro Gln Glu Gln Ala Thr Asp Pro Leu Val
690 695 700

Asp Ser Leu Gly Ser Gly Ile Val Tyr Ser Ala Leu Thr Cys His Leu
705 710 715 720

Cys Gly His Leu Lys Gln Cys His Gly Gln Glu Asp Gly Gly Gln Thr
725 730 735

Pro Val Met Ala Ser Pro Cys Cys Gly Cys Cys Cys Gly Asp Arg Ser
740 745 750

Ser Pro Pro Thr Thr Pro Leu Arg Ala Pro Asp Pro Ser Pro Gly Gly
755 760 765

Val Pro Leu Glu Ala Ser Leu Cys Pro Ala Ser Leu Ala Pro Ser Gly
770 775 780

Ile Ser Glu Lys Ser Lys Ser Ser Ser Ser Phe His Pro Ala Pro Gly
785 790 795 800

Asn Ala Gln Ser Ser Ser Gln Thr Pro Lys Ile Val Asn Phe Val Ser
805 810 815

Val Gly Pro Thr Tyr Met Arg Val Ser
820 825

<210> 30
<211> 9
<212> PRT
<213> Artificial

<220>
<223> Peptide

<400> 30

Phe Val Phe Ala Arg Thr Met Pro Ala
1 5

<210> 31
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 31

Asn Gly Pro Lys Ile Pro Ser Ile Ala Thr
1 5 10

<210> 32
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 32

Ala Thr Gly Gln Val Cys His Ala Leu
1 5

<210> 33

<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 33

Arg Lys Val Cys Asn Gly Ile Gly Ile Gly Glu
1 5 10

<210> 34
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 34

Trp His Asn Ser Tyr Arg Glu Pro Phe
1 5

<210> 35
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 35

Tyr Arg Glu Pro Phe Glu Gln His Leu Leu
1 5 10

<210> 36
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 36

Ser Asp Thr Leu Leu Thr Trp Ser
1 5

<210> 37
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 37

Ile Tyr Asn Val Thr Tyr Leu Glu
1 5

<210> 38
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 38

Ile Ala Ala Ser Thr Leu Lys Ser Gly Ile Ser
1 5 10

<210> 39
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 39

Lys Pro Ser Glu His Val Lys Pro Arg
1 5

<210> 40
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 40

Phe Thr Cys Glu Glu Asp Phe Tyr Phe Pro Trp
1 5 10

<210> 41
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 41

Ser Val Asp Glu Ile Val Gln Pro Asp
1 5

<210> 42
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 42

Met Asp Pro Ile Asp Thr Thr Ser Val Pro Val Tyr
1 5 10

<210> 43
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 43

Ile Asp Ala Ala Tyr Ile Gln Leu Ile Tyr Pro Val
1 5 10

<210> 44
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

Peptide

<400> 44

Leu Ile Tyr Pro Val Thr Asn Phe Gln Lys His Met
1 5 10

<210> 45

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 45

Leu Glu Glu Asn Lys Pro Thr Arg Pro Val
1 5 10

<210> 46

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 46

Asn Lys Pro Thr Arg Pro Val Ile Val Ser
1 5 10

<210> 47

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 47

Val Ala Glu Lys His Arg Gly Asn Tyr Thr
1 5 10

<210> 48

<211> 9

<212> PRT

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 48

Trp Asn Gly Ser Val Ile Asp Glu Asp
1 5

<210> 49
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 49

Val Pro Ala Pro Arg Tyr Thr Val Glu Leu
1 5 10

<210> 50
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 50

Ala Pro Arg Tyr Thr Val Glu Leu Ala
1 5

<210> 51
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 51

Val Gln Lys Asp Ser Cys Phe Asn Ser Pro Met
1 5 10

<210> 52

<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 52

Met Leu Pro Val His Lys Leu Tyr
1 5

<210> 53
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 53

Val Gly Ser Pro Lys Asn Ala Val Pro Pro Val
1 5 10

<210> 54
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 54

Val Thr Tyr Pro Glu Asn Gly Arg Thr Phe
1 5 10

<210> 55
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 55

Ile His Ser Pro Asn Asp His Val Val Tyr
1 5 10

<210> 56
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 56

Leu Ile Ser Asn Asn Gly Asn Tyr Thr
1 5

<210> 57
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 57

Val Trp Trp Thr Ile Asp Gly Lys Lys Pro Asp
1 5 10

<210> 58
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 58

Trp Thr Ile Asp Gly Lys Lys Pro Asp Asp Ile
1 5 10

<210> 59
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic Peptide

<400> 59

His Ser Arg Thr Glu Asp Glu Thr Arg Thr Gln
1 5 10

<210> 60
<211> 810
<212> PRT
<213> Mus sp.

<400> 60

Met Gly Arg Leu Cys Thr Lys Phe Leu Thr Ser Val Gly Cys Leu Ile
1 5 10 15

Leu Leu Leu Val Thr Gly Ser Gly Ser Ile Lys Val Leu Gly Glu Pro
20 25 30

Thr Cys Phe Ser Asp Tyr Ile Arg Thr Ser Thr Cys Glu Trp Phe Leu
35 40 45

Asp Ser Ala Val Asp Cys Ser Ser Gln Leu Cys Leu His Tyr Arg Leu
50 55 60

Met Phe Phe Glu Phe Ser Glu Asn Leu Thr Cys Ile Pro Arg Asn Ser
65 70 75 80

Ala Ser Thr Val Cys Val Cys His Met Glu Met Asn Arg Pro Val Gln
85 90 95

Ser Asp Arg Tyr Gln Met Glu Leu Trp Ala Glu His Arg Gln Leu Trp
100 105 110

Gln Gly Ser Phe Ser Pro Ser Gly Asn Val Lys Pro Leu Ala Pro Asp
115 120 125

Asn Leu Thr Leu His Thr Asn Val Ser Asp Glu Trp Leu Leu Thr Trp
130 135 140

Asn Asn Leu Tyr Pro Ser Asn Asn Leu Leu Tyr Lys Asp Leu Ile Ser
145 150 155 160

Met Val Asn Ile Ser Arg Glu Asp Asn Pro Ala Glu Phe Ile Val Tyr
165 170 175

Asn Val Thr Tyr Lys Glu Pro Arg Leu Ser Phe Pro Ile Asn Ile Leu

180	185	190
Met Ser Gly Val Tyr Tyr Thr Ala Arg Val Arg Val Arg Ser Gln Ile 195 200 205		
Leu Thr Gly Thr Trp Ser Glu Trp Ser Pro Ser Ile Thr Trp Tyr Asn 210 215 220		
His Phe Gln Leu Pro Leu Ile Gln Arg Leu Pro Leu Gly Val Thr Ile 225 230 235 240		
Ser Cys Leu Cys Ile Pro Leu Phe Cys Leu Phe Cys Tyr Phe Ser Ile 245 250 255		
Thr Lys Ile Lys Lys Ile Trp Trp Asp Gln Ile Pro Thr Pro Ala Arg 260 265 270		
Ser Pro Leu Val Ala Ile Ile Ile Gln Asp Ala Gln Val Pro Leu Trp 275 280 285		
Asp Lys Gln Thr Arg Ser Gln Glu Ser Thr Lys Tyr Pro His Trp Lys 290 295 300		
Thr Cys Leu Asp Lys Leu Leu Pro Cys Leu Leu Lys His Arg Val Lys 305 310 315 320		
Lys Lys Thr Asp Phe Pro Lys Ala Ala Pro Thr Lys Ser Leu Gln Ser 325 330 335		
Pro Gly Lys Ala Gly Trp Cys Pro Met Glu Val Ser Arg Thr Val Leu 340 345 350		
Trp Pro Glu Asn Val Ser Val Ser Val Val Arg Cys Met Glu Leu Phe 355 360 365		
Glu Ala Pro Val Gln Asn Val Glu Glu Glu Glu Asp Glu Ile Val Lys 370 375 380		
Glu Asp Leu Ser Met Ser Pro Glu Asn Ser Gly Gly Cys Gly Phe Gln 385 390 395 400		
Glu Ser Gln Ala Asp Ile Met Ala Arg Leu Thr Glu Asn Leu Phe Ser 405 410 415		

Asp Leu Leu Glu Ala Glu Asn Gly Gly Leu Gly Gln Ser Ala Leu Ala
420 425 430

Glu Ser Cys Ser Pro Leu Pro Ser Gly Ser Gly Gln Ala Ser Val Ser
435 440 445

Trp Ala Cys Leu Pro Met Gly Pro Ser Glu Glu Ala Thr Cys Gln Val
450 455 460

Thr Glu Gln Pro Ser His Pro Gly Pro Leu Ser Gly Ser Pro Ala Gln
465 470 475 480

Ser Ala Pro Thr Leu Ala Cys Thr Gln Val Pro Leu Val Leu Ala Asp
485 490 495

Asn Pro Ala Tyr Arg Ser Phe Ser Asp Cys Cys Ser Pro Ala Pro Asn
500 505 510

Pro Gly Glu Leu Ala Pro Glu Gln Gln Gln Ala Asp His Leu Glu Glu
515 520 525

Glu Glu Pro Pro Ser Pro Ala Asp Pro His Ser Ser Gly Pro Pro Met
530 535 540

Gln Pro Val Glu Ser Trp Glu Gln Ile Leu His Met Ser Val Leu Gln
545 550 555 560

His Gly Ala Ala Ala Gly Ser Thr Pro Ala Pro Ala Gly Gly Tyr Gln
565 570 575

Glu Phe Val Gln Ala Val Lys Gln Gly Ala Ala Gln Asp Pro Gly Val
580 585 590

Pro Gly Val Arg Pro Ser Gly Asp Pro Gly Tyr Lys Ala Phe Ser Ser
595 600 605

Leu Leu Ser Ser Asn Gly Ile Arg Gly Asp Thr Ala Ala Ala Gly Thr
610 615 620

Asp Asp Gly His Gly Gly Tyr Lys Pro Phe Gln Asn Pro Val Pro Asn
625 630 635 640

Gln Ser Pro Ser Ser Val Pro Leu Phe Thr Phe Gly Leu Asp Thr Glu
645 650 655

Leu Ser Pro Ser Pro Leu Asn Ser Asp Pro Pro Lys Ser Pro Pro Glu
660 665 670

Cys Leu Gly Leu Glu Leu Gly Leu Lys Gly Gly Asp Trp Val Lys Ala
675 680 685

Pro Pro Pro Ala Asp Gln Val Pro Lys Pro Phe Gly Asp Asp Leu Gly
690 695 700

Phe Gly Ile Val Tyr Ser Ser Leu Thr Cys His Leu Cys Gly His Leu
705 710 715 720

Lys Gln His His Ser Gln Glu Glu Gly Gly Gln Ser Pro Ile Val Ala
725 730 735

Ser Pro Gly Cys Gly Cys Cys Tyr Asp Asp Arg Ser Pro Ser Leu Gly
740 745 750

Ser Leu Ser Gly Ala Leu Glu Ser Cys Pro Glu Gly Ile Pro Pro Glu
755 760 765

Ala Asn Leu Met Ser Ala Pro Lys Thr Pro Ser Asn Leu Ser Gly Glu
770 775 780

Gly Lys Gly Pro Gly His Ser Pro Val Pro Ser Gln Thr Thr Glu Val
785 790 795 800

Pro Val Gly Ala Leu Gly Ile Ala Val Ser
805 810

<210> 61
<211> 810
<212> PRT
<213> Equus sp.

<400> 61

Met Gly Arg Leu Cys Thr Lys Phe Leu Thr Ser Val Gly Cys Leu Ile
1 5 10 15

Leu Leu Leu Val Thr Gly Ser Gly Ser Ile Lys Val Leu Gly Glu Pro

Copied from 10693357 on 07/15/2004

Thr Lys Ile Lys Lys Ile Trp Trp Asp Gln Ile Pro Thr Pro Ala Arg
260 265 270

Ser Pro Leu Val Ala Ile Ile Ile Gln Asp Ala Gln Val Pro Leu Trp
275 280 285

Asp Lys Gln Thr Arg Ser Gln Glu Ser Thr Lys Tyr Pro His Trp Lys
290 295 300

Thr Cys Leu Asp Lys Leu Leu Pro Cys Leu Leu Lys His Arg Val Lys
305 310 315 320

Lys Lys Thr Asp Phe Pro Lys Ala Ala Pro Thr Lys Ser Pro Gln Ser
325 330 335

Pro Gly Lys Ala Gly Trp Cys Pro Met Glu Val Ser Arg Thr Val Leu
340 345 350

Trp Pro Glu Asn Val Ser Val Ser Val Val Arg Cys Met Glu Leu Phe
355 360 365

Glu Ala Pro Val Gln Asn Val Glu Glu Glu Glu Asp Glu Ile Val Lys
370 375 380

Glu Asp Leu Ser Met Ser Pro Glu Asn Ser Gly Gly Cys Gly Phe Gln
385 390 395 400

Glu Ser Gln Ala Asp Ile Met Ala Arg Leu Thr Glu Asn Leu Phe Ser
405 410 415

Asp Leu Leu Glu Ala Glu Asn Gly Gly Leu Gly Gln Ser Ala Leu Ala
420 425 430

Glu Ser Cys Ser Pro Leu Pro Ser Gly Ser Gly Gln Ala Ser Val Ser
435 440 445

Trp Ala Cys Leu Pro Met Gly Pro Ser Glu Glu Ala Thr Cys Gln Val
450 455 460

Thr Glu Gln Pro Ser His Pro Gly Pro Leu Ser Gly Ser Pro Ala Gln
465 470 475 480

Ser Ala Pro Thr Leu Ala Cys Thr Gln Val Pro Leu Val Leu Ala Asp
485 490 495

Asn Pro Ala Tyr Arg Ser Phe Ser Asp Cys Cys Ser Pro Ala Pro Asn
500 505 510

Pro Gly Glu Leu Ala Pro Glu Gln Gln Gln Ala Asp His Leu Glu Glu
515 520 525

Glu Glu Pro Pro Ser Pro Ala Asp Pro His Ser Ser Gly Pro Pro Met
530 535 540

Gln Pro Val Glu Ser Trp Glu Gln Ile Leu His Met Ser Val Leu Gln
545 550 555 560

His Gly Ala Ala Ala Gly Ser Thr Pro Ala Pro Ala Gly Gly Tyr Gln
565 570 575

Glu Phe Val Gln Ala Val Lys Gln Gly Ala Ala Gln Asp Pro Gly Val
580 585 590

Pro Gly Val Arg Pro Ser Gly Asp Pro Gly Tyr Lys Ala Phe Ser Ser
595 600 605

Leu Leu Ser Ser Asn Gly Ile Arg Gly Asp Thr Ala Ala Ala Gly Thr
610 615 620

Asp Asp Gly His Gly Gly Tyr Lys Pro Phe Gln Asn Pro Val Pro Asn
625 630 635 640

Gln Ser Pro Ser Ser Val Pro Leu Phe Thr Phe Gly Leu Asp Thr Glu
645 650 655

Leu Ser Pro Ser Pro Leu Asn Ser Asp Pro Pro Lys Ser Pro Pro Glu
660 665 670

Cys Leu Gly Leu Glu Leu Gly Leu Lys Gly Gly Asp Trp Val Lys Ala
675 680 685

Pro Pro Pro Ala Asp Gln Val Pro Lys Pro Phe Gly Asp Asp Leu Gly
690 695 700

Phe Gly Ile Val Tyr Ser Ser Leu Thr Cys His Leu Cys Gly His Leu
705 710 715 720

Lys Gln His His Ser Gln Glu Glu Gly Gly Gln Ser Pro Ile Val Ala
725 730 735

Ser Pro Gly Cys Gly Cys Cys Tyr Asp Asp Arg Ser Pro Ser Leu Gly
740 745 750

Ser Leu Ser Gly Ala Leu Glu Ser Cys Pro Glu Gly Ile Pro Pro Glu
755 760 765

Ala Asn Leu Met Ser Ala Pro Lys Thr Pro Ser Asn Leu Ser Gly Glu
770 775 780

Gly Lys Gly Pro Gly His Ser Pro Val Pro Ser Gln Thr Thr Glu Val
785 790 795 800

Pro Val Gly Ala Leu Gly Ile Ala Val Ser
805 810

<210> 62

<211> 1356

<212> PRT

<213> Homo sapiens

<400> 62

Met Gln Ser Lys Val Leu Leu Ala Val Ala Leu Trp Leu Cys Val Glu
1 5 10 15

Thr Arg Ala Ala Ser Val Gly Leu Pro Ser Val Ser Leu Asp Leu Pro
20 25 30

Arg Leu Ser Ile Gln Lys Asp Ile Leu Thr Ile Lys Ala Asn Thr Thr
35 40 45

Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro
50 55 60

Asn Asn Gln Ser Gly Ser Glu Gln Arg Val Glu Val Thr Glu Cys Ser
65 70 75 80

Asp Gly Leu Phe Cys Lys Thr Leu Thr Ile Pro Lys Val Ile Gly Asn
85 90 95

Asp Thr Gly Ala Tyr Lys Cys Phe Tyr Arg Glu Thr Asp Leu Ala Ser
 100 105 110

Val Ile Tyr Val Tyr Val Gln Asp Tyr Arg Ser Pro Phe Ile Ala Ser
 115 120 125

Val Ser Asp Gln His Gly Val Val Tyr Ile Thr Glu Asn Lys Asn Lys
 130 135 140

Thr Val Val Ile Pro Cys Leu Gly Ser Ile Ser Asn Leu Asn Val Ser
 145 150 155 160

Leu Cys Ala Arg Tyr Pro Glu Lys Arg Phe Val Pro Asp Gly Asn Arg
 165 170 175

Ile Ser Trp Asp Ser Lys Lys Gly Phe Thr Ile Pro Ser Tyr Met Ile
 180 185 190

Ser Tyr Ala Gly Met Val Phe Cys Glu Ala Lys Ile Asn Asp Glu Ser
 195 200 205

Tyr Gln Ser Ile Met Tyr Ile Val Val Val Val Gly Tyr Arg Ile Tyr
 210 215 220

Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu
 225 230 235 240

Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile
 245 250 255

Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu
 260 265 270

Val Asn Arg Asp Leu Lys Thr Gln Ser Gly Ser Glu Met Lys Lys Phe
 275 280 285

Leu Ser Thr Leu Thr Ile Asp Gly Val Thr Arg Ser Asp Gln Gly Leu
 290 295 300

Tyr Thr Cys Ala Ala Ser Ser Gly Leu Met Thr Lys Lys Asn Ser Thr
 305 310 315 320

Phe Val Arg Val His Glu Lys Pro Phe Val Ala Phe Gly Ser Gly Met
325 330 335

Glu Ser Leu Val Glu Ala Thr Val Gly Glu Arg Val Arg Ile Pro Ala
340 345 350

Lys Tyr Leu Gly Tyr Pro Pro Pro Glu Ile Lys Trp Tyr Lys Asn Gly
355 360 365

Ile Pro Leu Glu Ser Asn His Thr Ile Lys Ala Gly His Val Leu Thr
370 375 380

Ile Met Glu Val Ser Glu Arg Asp Thr Gly Asn Tyr Thr Val Ile Leu
385 390 395 400

Thr Asn Pro Ile Ser Lys Glu Lys Gln Ser His Val Val Ser Leu Val
405 410 415

Val Tyr Val Pro Pro Gln Ile Gly Glu Lys Ser Leu Ile Ser Pro Val
420 425 430

Asp Ser Tyr Gln Tyr Gly Thr Thr Gln Thr Leu Thr Cys Thr Val Tyr
435 440 445

Ala Ile Pro Pro Pro His His Ile His Trp Tyr Trp Gln Leu Glu Glu
450 455 460

Glu Cys Ala Asn Glu Pro Ser Gln Ala Val Ser Val Thr Asn Pro Tyr
465 470 475 480

Pro Cys Glu Glu Trp Arg Ser Val Glu Asp Phe Gln Gly Gly Asn Lys
485 490 495

Ile Glu Val Asn Lys Asn Gln Phe Ala Leu Ile Glu Gly Lys Asn Lys
500 505 510

Thr Val Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr
515 520 525

Lys Cys Glu Ala Val Asn Lys Val Gly Arg Gly Glu Arg Val Ile Ser
530 535 540

Phe His Val Thr Arg Gly Pro Glu Ile Thr Leu Gln Pro Asp Met Gln
545 550 555 560

Pro Thr Glu Gln Glu Ser Val Ser Leu Trp Cys Thr Ala Asp Arg Ser
565 570 575

Thr Phe Glu Asn Leu Thr Trp Tyr Lys Leu Gly Pro Gln Pro Leu Pro
580 585 590

Ile His Val Gly Glu Leu Pro Thr Pro Val Cys Lys Asn Leu Asp Thr
595 600 605

Leu Trp Lys Leu Asn Ala Thr Met Phe Ser Asn Ser Thr Asn Asp Ile
610 615 620

Leu Ile Met Glu Leu Lys Asn Ala Ser Leu Gln Asp Gln Gly Asp Tyr
625 630 635 640

Val Cys Leu Ala Gln Asp Arg Lys Thr Lys Lys Arg His Cys Val Val
645 650 655

Arg Gln Leu Thr Val Leu Glu Arg Val Ala Pro Thr Ile Thr Gly Asn
660 665 670

Leu Glu Asn Gln Thr Thr Ser Ile Gly Glu Ser Ile Glu Val Ser Cys
675 680 685

Thr Ala Ser Gly Asn Pro Pro Pro Gln Ile Met Trp Phe Lys Asp Asn
690 695 700

Glu Thr Leu Val Glu Asp Ser Gly Ile Val Leu Lys Asp Gly Asn Arg
705 710 715 720

Asn Leu Thr Ile Arg Arg Val Arg Lys Glu Asp Glu Gly Leu Tyr Thr
725 730 735

Cys Gln Ala Cys Ser Val Leu Gly Cys Ala Lys Val Glu Ala Phe Phe
740 745 750

Ile Ile Glu Gly Ala Gln Glu Lys Thr Asn Leu Glu Ile Ile Ile Leu
755 760 765

Val Gly Thr Ala Val Ile Ala Met Phe Phe Trp Leu Leu Leu Val Ile

770	775	780
Ile Leu Arg Thr Val Lys Arg Ala Asn Gly Gly Glu Leu Lys Thr Gly 785 790 795 800		
Tyr Leu Ser Ile Val Met Asp Pro Asp Glu Leu Pro Leu Asp Glu His 805 810 815		
Cys Glu Arg Leu Pro Tyr Asp Ala Ser Lys Trp Glu Phe Pro Arg Asp 820 825 830		
Arg Leu Lys Leu Gly Lys Pro Leu Gly Arg Gly Ala Phe Gly Gln Val 835 840 845		
Ile Glu Ala Asp Ala Phe Gly Ile Asp Lys Thr Ala Thr Cys Arg Thr 850 855 860		
Val Ala Val Lys Met Leu Lys Glu Gly Ala Thr His Ser Glu His Arg 865 870 875 880		
Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly His His Leu 885 890 895		
Asn Val Val Asn Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu 900 905 910		
Met Val Ile Val Glu Phe Cys Lys Phe Gly Asn Leu Ser Thr Tyr Leu 915 920 925		
Arg Ser Lys Arg Asn Glu Phe Val Pro Tyr Lys Thr Lys Gly Ala Arg 930 935 940		
Phe Arg Gln Gly Lys Asp Tyr Val Gly Ala Ile Pro Val Asp Leu Lys 945 950 955 960		
Arg Arg Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly 965 970 975		
Phe Val Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Pro 980 985 990		
Glu Asp Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr 995 1000 1005		

Ser	Phe	Gln	Val	Ala	Lys	Gly	Met	Glu	Phe	Leu	Ala	Ser	Arg	Lys
1010						1015					1020			
Cys	Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Leu	Ser	Glu
1025						1030					1035			
Lys	Asn	Val	Val	Lys	Ile	Cys	Asp	Phe	Gly	Leu	Ala	Arg	Asp	Ile
1040						1045					1050			
Tyr	Lys	Asp	Pro	Asp	Tyr	Val	Arg	Lys	Gly	Asp	Ala	Arg	Leu	Pro
1055						1060					1065			
Leu	Lys	Trp	Met	Ala	Pro	Glu	Thr	Ile	Phe	Asp	Arg	Val	Tyr	Thr
1070						1075					1080			
Ile	Gln	Ser	Asp	Val	Trp	Ser	Phe	Gly	Val	Leu	Leu	Trp	Glu	Ile
1085						1090					1095			
Phe	Ser	Leu	Gly	Ala	Ser	Pro	Tyr	Pro	Gly	Val	Lys	Ile	Asp	Glu
1100						1105					1110			
Glu	Phe	Cys	Arg	Arg	Leu	Lys	Glu	Gly	Thr	Arg	Met	Arg	Ala	Pro
1115						1120					1125			
Asp	Tyr	Thr	Thr	Pro	Glu	Met	Tyr	Gln	Thr	Met	Leu	Asp	Cys	Trp
1130						1135					1140			
His	Gly	Glu	Pro	Ser	Gln	Arg	Pro	Thr	Phe	Ser	Glu	Leu	Val	Glu
1145						1150					1155			
His	Leu	Gly	Asn	Leu	Leu	Gln	Ala	Asn	Ala	Gln	Gln	Asp	Gly	Lys
1160						1165					1170			
Asp	Tyr	Ile	Val	Leu	Pro	Ile	Ser	Glu	Thr	Leu	Ser	Met	Glu	Glu
1175						1180					1185			
Asp	Ser	Gly	Leu	Ser	Leu	Pro	Thr	Ser	Pro	Val	Ser	Cys	Met	Glu
1190						1195					1200			
Glu	Glu	Glu	Val	Cys	Asp	Pro	Lys	Phe	His	Tyr	Asp	Asn	Thr	Ala
1205						1210					1215			

Gly Ile Ser Gln Tyr Leu Gln Asn Ser Lys Arg Lys Ser Arg Pro
1220 1225 1230

Val Ser Val Lys Thr Phe Glu Asp Ile Pro Leu Glu Glu Pro Glu
1235 1240 1245

Val Lys Val Ile Pro Asp Asp Asn Gln Thr Asp Ser Gly Met Val
1250 1255 1260

Leu Ala Ser Glu Glu Leu Lys Thr Leu Glu Asp Arg Thr Lys Leu
1265 1270 1275

Ser Pro Ser Phe Gly Gly Met Val Pro Ser Lys Ser Arg Glu Ser
1280 1285 1290

Val Ala Ser Glu Gly Ser Asn Gln Thr Ser Gly Tyr Gln Ser Gly
1295 1300 1305

Tyr His Ser Asp Asp Thr Asp Thr Thr Val Tyr Ser Ser Glu Glu
1310 1315 1320

Ala Glu Leu Leu Lys Leu Ile Glu Ile Gly Val Gln Thr Gly Ser
1325 1330 1335

Thr Ala Gln Ile Leu Gln Pro Asp Ser Gly Thr Thr Leu Ser Ser
1340 1345 1350

Pro Pro Val
1355

<210> 63
<211> 1367
<212> PRT
<213> Mus sp.

<400> 63

Met Glu Ser Lys Ala Leu Leu Ala Val Ala Leu Trp Phe Cys Val Glu
1 5 10 15

Thr Arg Ala Ala Ser Val Gly Leu Pro Gly Asp Phe Leu His Pro Pro
20 25 30

Lys Leu Ser Thr Gln Lys Asp Ile Leu Thr Ile Leu Ala Asn Thr Thr

35

40

45

Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro
50 55 60

Asn Ala Gln Arg Asp Ser Glu Glu Arg Val Leu Val Thr Glu Cys Gly
65 70 75 80

Gly Gly Asp Ser Ile Phe Cys Lys Thr Leu Thr Ile Pro Arg Val Val
85 90 95

Gly Asn Asp Thr Gly Ala Tyr Lys Cys Ser Tyr Arg Asp Val Asp Ile
100 105 110

Ala Ser Thr Val Tyr Val Tyr Val Arg Asp Tyr Arg Ser Pro Phe Ile
115 120 125

Ala Ser Val Ser Asp Gln His Gly Ile Val Tyr Ile Thr Glu Asn Lys
130 135 140

Asn Lys Thr Val Val Ile Pro Cys Arg Gly Ser Ile Ser Asn Leu Asn
145 150 155 160

Val Ser Leu Cys Ala Arg Tyr Pro Glu Lys Arg Phe Val Pro Asp Gly
165 170 175

Asn Arg Ile Ser Trp Asp Ser Glu Ile Gly Phe Thr Leu Pro Ser Tyr
180 185 190

Met Ile Ser Tyr Ala Gly Met Val Phe Cys Glu Ala Lys Ile Asn Asp
195 200 205

Glu Thr Tyr Gln Ser Ile Met Tyr Ile Val Val Val Gly Tyr Arg
210 215 220

Ile Tyr Asp Val Ile Leu Ser Pro Pro His Glu Ile Glu Leu Ser Ala
225 230 235 240

Gly Glu Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val
245 250 255

Gly Leu Asp Phe Thr Trp His Ser Pro Pro Ser Lys Ser His His Lys
260 265 270

Lys Ile Val Asn Arg Asp Val Lys Pro Phe Pro Gly Thr Val Ala Lys
 275 280 285

Met Phe Leu Ser Thr Leu Thr Ile Glu Ser Val Thr Lys Ser Asp Gln
 290 295 300

Gly Glu Tyr Thr Cys Val Ala Ser Ser Gly Arg Met Ile Lys Arg Asn
 305 310 315 320

Arg Thr Phe Val Arg Val His Thr Lys Pro Phe Ile Ala Phe Gly Ser
 325 330 335

Gly Met Lys Ser Leu Val Glu Ala Thr Val Gly Ser Gln Val Arg Ile
 340 345 350

Pro Val Lys Tyr Leu Ser Tyr Pro Ala Pro Asp Ile Lys Trp Tyr Arg
 355 360 365

Asn Gly Arg Pro Ile Glu Ser Asn Tyr Thr Met Ile Val Gly Asp Glu
 370 375 380

Leu Thr Ile Met Glu Val Thr Glu Arg Asp Ala Gly Asn Tyr Thr Val
 385 390 395 400

Ile Leu Thr Asn Pro Ile Ser Met Glu Lys Gln Ser His Met Val Ser
 405 410 415

Leu Val Val Asn Val Pro Pro Gln Ile Gly Glu Lys Ala Leu Ile Ser
 420 425 430

Pro Met Asp Ser Tyr Gln Tyr Gly Thr Met Gln Thr Leu Thr Cys Thr
 435 440 445

Val Tyr Ala Asn Pro Pro Leu His His Ile Gln Trp Tyr Trp Gln Leu
 450 455 460

Glu Glu Ala Cys Ser Tyr Arg Pro Gly Gln Thr Ser Pro Tyr Ala Cys
 465 470 475 480

Lys Glu Trp Arg His Val Glu Asp Phe Gln Gly Gly Asn Lys Ile Glu
 485 490 495

Val Thr Lys Asn Gln Tyr Ala Leu Ile Glu Gly Lys Asn Lys Thr Val
500 505 510

Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr Lys Cys
515 520 525

Glu Ala Ile Asn Lys Ala Gly Arg Gly Glu Arg Val Ile Ser Phe His
530 535 540

Val Ile Arg Gly Pro Glu Ile Thr Val Gln Pro Ala Ala Gln Pro Thr
545 550 555 560

Glu Gln Glu Ser Val Ser Leu Leu Cys Thr Ala Asp Arg Asn Thr Phe
565 570 575

Glu Asn Leu Thr Trp Tyr Lys Leu Gly Ser Gln Ala Thr Ser Val His
580 585 590

Met Gly Glu Ser Leu Thr Pro Val Cys Lys Asn Leu Asp Ala Leu Trp
595 600 605

Lys Leu Asn Gly Thr Met Phe Ser Asn Ser Thr Asn Asp Ile Leu Ile
610 615 620

Val Ala Phe Gln Asn Ala Ser Leu Gln Asp Gln Gly Asp Tyr Val Cys
625 630 635 640

Ser Ala Gln Asp Lys Lys Thr Lys Lys Arg His Cys Leu Val Lys Gln
645 650 655

Leu Ile Ile Leu Glu Arg Met Ala Pro Met Ile Thr Gly Asn Leu Glu
660 665 670

Asn Gln Thr Thr Thr Ile Gly Glu Thr Ile Glu Val Thr Cys Pro Ala
675 680 685

Ser Gly Asn Pro Thr Pro His Ile Thr Trp Phe Lys Asp Asn Glu Thr
690 695 700

Leu Val Glu Asp Ser Gly Ile Val Leu Arg Asp Gly Asn Arg Asn Leu
705 710 715 720

Thr Ile Arg Arg Val Arg Lys Glu Asp Gly Gly Leu Tyr Thr Cys Gln
725 730 735

Ala Cys Asn Val Leu Gly Cys Ala Arg Ala Glu Thr Leu Phe Ile Ile
740 745 750

Glu Gly Ala Gln Glu Lys Thr Asn Leu Glu Val Ile Ile Leu Val Gly
755 760 765

Thr Ala Val Ile Ala Met Phe Phe Trp Leu Leu Leu Val Ile Leu Val
770 775 780

Arg Thr Val Lys Arg Ala Asn Glu Gly Glu Leu Lys Thr Gly Tyr Leu
785 790 795 800

Ser Ile Val Met Asp Pro Asp Glu Leu Pro Leu Asp Glu Arg Cys Glu
805 810 815

Arg Leu Pro Tyr Asp Ala Ser Lys Trp Glu Phe Pro Arg Asp Arg Leu
820 825 830

Lys Leu Gly Lys Pro Leu Gly Arg Gly Ala Phe Gly Gln Val Ile Glu
835 840 845

Ala Asp Ala Phe Gly Ile Asp Lys Thr Ala Thr Cys Lys Thr Val Ala
850 855 860

Val Lys Met Leu Lys Glu Gly Ala Thr His Ser Glu His Arg Ala Leu
865 870 875 880

Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly His His Leu Asn Val
885 890 895

Val Asn Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu Met Val
900 905 910

Ile Val Glu Phe Ser Lys Phe Gly Asn Leu Ser Thr Tyr Leu Arg Gly
915 920 925

Lys Arg Asn Glu Phe Val Pro Tyr Lys Ser Lys Gly Ala Arg Phe Arg
930 935 940

Gln Gly Lys Asp Tyr Val Gly Glu Leu Ser Val Asp Leu Lys Arg Arg

945	950	955	960
Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val			
	965	970	975
Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Ser Glu Glu			
	980	985	990
Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe			
	995	1000	1005
Gln Val Ala Lys Gly Met Glu Phe Leu Ala Ser Arg Lys Cys Ile			
	1010	1015	1020
His Arg Asp Leu Ala Ala Arg Asn Ile Leu Leu Ser Glu Lys Asn			
	1025	1030	1035
Val Val Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Tyr Lys			
	1040	1045	1050
Asp Pro Asp Tyr Val Arg Lys Gly Asp Ala Arg Leu Pro Leu Lys			
	1055	1060	1065
Trp Met Ala Pro Glu Thr Ile Phe Asp Arg Val Tyr Thr Ile Gln			
	1070	1075	1080
Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe Ser			
	1085	1090	1095
Leu Gly Ala Ser Pro Tyr Pro Gly Val Lys Ile Asp Glu Glu Phe			
	1100	1105	1110
Cys Arg Arg Leu Lys Glu Gly Thr Arg Met Arg Ala Pro Asp Tyr			
	1115	1120	1125
Thr Thr Pro Glu Met Tyr Gln Thr Met Leu Asp Cys Trp His Glu			
	1130	1135	1140
Asp Pro Asn Gln Arg Pro Ser Phe Ser Glu Leu Val Glu His Leu			
	1145	1150	1155
Gly Asn Leu Leu Gln Ala Asn Ala Gln Gln Asp Gly Lys Asp Tyr			
	1160	1165	1170

Ile Val Leu Pro Met Ser Glu Thr Leu Ser Met Glu Glu Asp Ser
1175 1180 1185

Gly Leu Ser Leu Pro Thr Ser Pro Val Ser Cys Met Glu Glu Glu
1190 1195 1200

Glu Val Cys Asp Pro Lys Phe His Tyr Asp Asn Thr Ala Gly Ile
1205 1210 1215

Ser His Tyr Leu Gln Asn Ser Lys Arg Lys Ser Arg Pro Val Ser
1220 1225 1230

Val Lys Thr Phe Glu Asp Ile Pro Leu Glu Glu Pro Glu Val Lys
1235 1240 1245

Val Ile Pro Asp Asp Ser Gln Thr Asp Ser Gly Met Val Leu Ala
1250 1255 1260

Ser Glu Glu Leu Lys Thr Leu Glu Asp Arg Asn Lys Leu Ser Pro
1265 1270 1275

Ser Phe Gly Gly Met Met Pro Ser Lys Ser Arg Glu Ser Val Ala
1280 1285 1290

Ser Glu Gly Ser Asn Gln Thr Ser Gly Tyr Gln Ser Gly Tyr His
1295 1300 1305

Ser Asp Asp Thr Asp Thr Thr Val Tyr Ser Ser Asp Glu Ala Gly
1310 1315 1320

Leu Leu Lys Met Val Asp Ala Ala Val His Ala Asp Ser Gly Thr
1325 1330 1335

Thr Leu Gln Leu Thr Ser Cys Leu Asn Gly Ser Gly Pro Val Pro
1340 1345 1350

Ala Pro Pro Pro Thr Pro Gly Asn His Glu Arg Gly Ala Ala
1355 1360 1365

<210> 64
<211> 1343
<212> PRT

<213> Rattus sp.

<400> 64

Met Glu Ser Arg Ala Leu Leu Ala Val Ala Leu Trp Phe Cys Val Glu
1 5 10 15

Thr Arg Ala Ala Ser Val Gly Leu Pro Gly Asp Ser Leu His Pro Pro
20 25 30

Lys Leu Ser Thr Gln Lys Asp Ile Leu Thr Ile Leu Ala Asn Thr Thr
35 40 45

Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro
50 55 60

Asn Thr Pro Arg Asp Ser Glu Glu Arg Val Leu Val Thr Glu Cys Gly
65 70 75 80

Asp Ser Ile Phe Cys Lys Thr Leu Thr Val Pro Arg Val Val Gly Asn
85 90 95

Asp Thr Gly Ala Tyr Lys Cys Phe Tyr Arg Asp Thr Asp Val Ser Ser
100 105 110

Ile Val Tyr Val Tyr Val Gln Asp His Arg Ser Pro Phe Ile Ala Ser
115 120 125

Val Ser Asp Glu His Gly Ile Val Tyr Ile Thr Glu Asn Lys Asn Lys
130 135 140

Thr Val Val Ile Pro Cys Arg Gly Ser Ile Ser Asn Leu Asn Val Ser
145 150 155 160

Leu Cys Ala Arg Tyr Pro Glu Lys Arg Phe Val Pro Asp Gly Asn Arg
165 170 175

Ile Ser Trp Asp Ser Glu Lys Gly Phe Thr Ile Pro Ser Tyr Met Ile
180 185 190

Ser Tyr Ala Gly Met Val Phe Cys Glu Ala Lys Ile Asn Asp Glu Thr
195 200 205

Tyr Gln Ser Ile Met Tyr Ile Val Leu Val Val Gly Tyr Arg Ile Tyr

210	215	220
Asp Val Val Leu Ser Pro Pro His Glu Ile Glu Leu Ser Ala Gly Glu 225 230 235 240		
Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Leu 245 250 255		
Asp Phe Ser Trp Gln Phe Pro Ser Ser Lys His Gln His Lys Lys Ile 260 265 270		
Val Asn Arg Asp Val Lys Ser Leu Pro Gly Thr Val Ala Lys Met Phe 275 280 285		
Leu Ser Thr Leu Thr Ile Asp Ser Val Thr Lys Ser Asp Gln Gly Glu 290 295 300		
Tyr Thr Cys Thr Ala Tyr Ser Gly Leu Met Thr Lys Lys Asn Lys Thr 305 310 315 320		
Phe Val Arg Val His Thr Lys Pro Phe Ile Ala Phe Gly Ser Gly Met 325 330 335		
Lys Ser Leu Val Glu Ala Thr Val Gly Ser Gln Val Arg Ile Pro Val 340 345 350		
Lys Tyr Leu Ser Tyr Pro Ala Pro Asp Ile Lys Trp Tyr Arg Asn Gly 355 360 365		
Arg Pro Ile Glu Ser Asn Tyr Thr Met Ile Val Gly Asp Glu Leu Thr 370 375 380		
Ile Met Glu Val Ser Glu Arg Asp Ala Gly Asn Tyr Thr Val Ile Leu 385 390 395 400		
Thr Asn Pro Ile Ser Met Glu Lys Gln Ser His Met Val Ser Leu Val 405 410 415		
Val Asn Val Pro Pro Gln Ile Gly Glu Lys Ala Leu Ile Ser Pro Met 420 425 430		
Asp Ser Tyr Gln Tyr Gly Thr Met Gln Thr Leu Thr Cys Thr Val Tyr 435 440 445		

Ala Asn Pro Pro Leu His His Ile Gln Trp Tyr Trp Gln Leu Glu Glu
450 455 460

Ala Cys Ser Tyr Arg Pro Ser Gln Thr Asn Pro Tyr Thr Cys Lys Glu
465 470 475 480

Trp Arg His Val Lys Asp Phe Gln Gly Gly Asn Lys Ile Glu Val Thr
485 490 495

Lys Asn Gln Tyr Ala Leu Ile Glu Gly Lys Asn Lys Thr Val Ser Thr
500 505 510

Leu Val Ile Gln Ala Ala Tyr Val Ser Ala Leu Tyr Lys Cys Glu Ala
515 520 525

Ile Asn Lys Ala Gly Arg Gly Glu Arg Val Ile Ser Phe His Val Ile
530 535 540

Arg Gly Pro Glu Ile Thr Val Gln Pro Ala Thr Gln Pro Thr Glu Arg
545 550 555 560

Glu Ser Met Ser Leu Leu Cys Thr Ala Asp Arg Asn Thr Phe Glu Asn
565 570 575

Leu Thr Trp Tyr Lys Leu Gly Ser Gln Ala Thr Ser Val His Met Gly
580 585 590

Glu Ser Leu Thr Pro Val Cys Lys Asn Leu Asp Ala Leu Trp Lys Leu
595 600 605

Asn Gly Thr Val Phe Ser Asn Ser Thr Asn Asp Ile Leu Ile Val Ala
610 615 620

Phe Gln Asn Ala Ser Leu Gln Asp Gln Gly Asn Tyr Val Cys Ser Ala
625 630 635 640

Gln Asp Lys Lys Thr Lys Lys Arg His Cys Leu Val Lys Gln Leu Val
645 650 655

Ile Leu Glu Arg Met Ala Pro Met Ile Thr Gly Asn Leu Glu Asn Gln
660 665 670

Thr Thr Thr Ile Gly Glu Thr Ile Glu Val Val Cys Pro Thr Ser Gly
675 680 685

Asn Pro Thr Pro Leu Ile Thr Trp Phe Lys Asp Asn Glu Thr Leu Val
690 695 700

Glu Asp Ser Gly Ile Val Leu Lys Asp Gly Asn Arg Asn Leu Thr Ile
705 710 715 720

Arg Arg Val Arg Lys Glu Asp Gly Gly Leu Tyr Thr Cys Gln Ala Cys
725 730 735

Asn Val Leu Gly Cys Ala Arg Ala Glu Thr Leu Phe Ile Ile Glu Gly
740 745 750

Val Gln Glu Lys Thr Asn Leu Glu Val Ile Ile Leu Val Gly Thr Ala
755 760 765

Val Ile Ala Met Phe Phe Trp Leu Leu Leu Val Ile Leu Val Arg Thr
770 775 780

Val Lys Arg Ala Asn Glu Gly Glu Leu Lys Thr Gly Tyr Leu Ser Ile
785 790 795 800

Val Met Asp Pro Asp Glu Leu Pro Leu Asp Glu Arg Cys Glu Arg Leu
805 810 815

Pro Tyr Asp Ala Ser Lys Trp Glu Phe Pro Arg Asp Arg Leu Lys Leu
820 825 830

Gly Lys Pro Leu Gly Arg Gly Ala Phe Gly Gln Val Ile Glu Ala Asp
835 840 845

Ala Phe Gly Ile Asp Lys Thr Ala Thr Cys Lys Thr Val Ala Val Lys
850 855 860

Met Leu Lys Glu Gly Ala Thr His Ser Glu His Arg Ala Leu Met Ser
865 870 875 880

Glu Leu Lys Ile Leu Ile His Ile Gly His His Leu Asn Val Val Asn
885 890 895

Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu Met Val Ile Val
900 905 910

Glu Phe Cys Lys Phe Gly Asn Leu Ser Thr Tyr Leu Arg Gly Lys Arg
915 920 925

Asn Glu Phe Val Pro Tyr Lys Ser Lys Gly Ala Arg Phe Arg Ser Gly
930 935 940

Lys Asp Tyr Val Gly Glu Leu Ser Val Asp Leu Lys Arg Arg Leu Asp
945 950 955 960

Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val Glu Glu
965 970 975

Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Ser Glu Glu Leu Tyr
980 985 990

Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe Gln Val
995 1000 1005

Ala Lys Gly Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg
1010 1015 1020

Asp Leu Ala Ala Arg Asn Ile Leu Leu Ser Glu Lys Asn Val Val
1025 1030 1035

Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro
1040 1045 1050

Asp Tyr Val Arg Lys Gly Asp Pro Arg Leu Pro Leu Lys Trp Met
1055 1060 1065

Ala Pro Glu Thr Ile Phe Asp Arg Ile Tyr Thr Ile Gln Ser Gly
1070 1075 1080

Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe Ser Leu Gly
1085 1090 1095

Ala Ser Pro Tyr Pro Gly Val Lys Ile Asp Glu Lys Phe Cys Arg
1100 1105 1110

Arg Leu Lys Glu Gly Thr Arg Met Arg Ala Pro Asp Tyr Thr Thr

1115		1120		1125
Pro Glu Met Tyr Gln Thr Met	Leu Asp Cys Trp His	Glu Asp Pro		
1130	1135	1140		
Asn Gln Arg Pro Ala Phe Ser	Glu Leu Val Glu His	Leu Gly Asn		
1145	1150	1155		
Leu Leu Gln Ala Asn Ala Gln	Gln Asp Gly Lys Asp	Tyr Ile Val		
1160	1165	1170		
Leu Pro Met Ser Glu Thr Leu	Ser Met Glu Glu Asp	Ser Gly Leu		
1175	1180	1185		
Ser Leu Pro Thr Ser Pro Val	Ser Cys Met Glu Glu	Glu Glu Val		
1190	1195	1200		
Cys Asp Pro Lys Phe His Tyr	Asp Asn Thr Ala Gly	Ile Ser His		
1205	1210	1215		
Tyr Leu Gln Asn Ser Lys Arg	Lys Ser Arg Pro Val	Ser Val Lys		
1220	1225	1230		
Thr Phe Glu Asp Ile Pro Leu	Glu Glu Pro Glu Val	Lys Val Ile		
1235	1240	1245		
Pro Asp Asp Ser Gln Thr Asp	Ser Gly Met Val Leu	Ala Ser Glu		
1250	1255	1260		
Glu Leu Lys Thr Leu Glu Asp	Arg Asn Lys Leu Ser	Pro Ser Phe		
1265	1270	1275		
Gly Gly Met Met Pro Ser Lys	Ser Arg Glu Ser Val	Ala Ser Glu		
1280	1285	1290		
Gly Ser Asn Gln Thr Ser Gly	Tyr Gln Ser Gly Tyr	His Ser Asp		
1295	1300	1305		
Asp Thr Asp Thr Thr Val Tyr	Ser Ser Asp Glu Ala	Gly Leu Leu		
1310	1315	1320		
Lys Leu Val Asp Val Ala Gly	His Val Asp Ser Gly	Thr Thr Leu		
1325	1330	1335		

Arg Ser Ser Pro Val
1340

<210> 65
<211> 1348
<212> PRT
<213> Callipepla sp.

<400> 65

Met Glu Leu Gly Pro Leu Arg Val Leu Thr Val Leu Leu Cys Leu Ala
1 5 10 15

Pro Val Phe Ala Gly Leu Phe Ile Ser Met Asp Gln Pro Thr Leu Ser
20 25 30

Ile Gln Lys Ser Val Leu Thr Ile Thr Thr Asn Asp Thr Leu Asn Ile
35 40 45

Thr Cys Ser Gly Gln Arg Ala Val Tyr Trp Ser Trp Pro Asn Asn Gln
50 55 60

Ser Ser Val Glu Lys Arg Leu Ala Val Thr Gly Cys Ser Glu Gly Pro
65 70 75 80

Phe Cys Lys Thr Leu Thr Leu Leu Arg Val Ile Gly Asn Asp Thr Gly
85 90 95

Asp Tyr Arg Cys Leu Tyr Gly Asp Ser Gln Ala Ala Thr Thr Ile Tyr
100 105 110

Val Tyr Val Gln Asp Tyr Arg Ser Pro Phe Val Thr Ser Val Gly Asp
115 120 125

Gln Leu Gly Ile Val Tyr Ile Thr Lys Asn Lys Thr Val Val Val Pro
130 135 140

Cys Leu Gly Thr Val Ser Asn Leu Asn Val Ser Leu His Ala Lys Tyr
145 150 155 160

Pro Glu Lys Val Phe Val Pro Asp Gly Lys Ser Ile Ser Trp Asp Asn
165 170 175

Lys Lys Gly Phe Thr Ile Pro Ser His Leu Ile Asn Tyr Ala Gly Met
180 185 190

Val Phe Cys Glu Ala Lys Ile Asp Asn Glu Ser Tyr Gln Ser Val Ile
195 200 205

Tyr Ile Val Ala Val Val Gly Tyr Arg Ile Tyr Asp Leu Thr Met Asn
210 215 220

Pro His Tyr Gln Val Glu Leu Ala Val Gly Glu Lys Leu Val Leu Asn
225 230 235 240

Cys Thr Val Arg Thr Glu Leu Asn Val Gly Ile Asp Phe Arg Trp Asp
245 250 255

Tyr Pro Ser Ile Lys Glu Arg Arg Ala Thr Ile Arg Asp Leu Lys Thr
260 265 270

Thr Ala Gly Glu Ile Lys Thr Phe Val Ser Thr Leu Thr Ile Glu Ser
275 280 285

Val Asn Leu Ser Asp Lys Gly Arg Tyr Thr Cys Ala Ala Ser Ser Gly
290 295 300

Arg Met Asn Met Lys Asn Ser Ser Tyr Phe Ile Ile His Glu Ser Pro
305 310 315 320

Phe Ile His Leu Glu Lys Met Glu Asn Val Val Glu Met Lys Leu Gly
325 330 335

Asp Thr Val Ser Ile Pro Val Lys Phe Lys Gly Tyr Pro Pro Pro Glu
340 345 350

Ala Lys Trp Tyr Lys Asn Gly Lys Val Ile Asn Ala Asn His Thr Val
355 360 365

Lys Leu Gly Tyr Ala Leu Val Ile Thr Glu Ala Thr Glu Lys Asp Ala
370 375 380

Gly Asn Tyr Thr Val Val Leu Thr Asn Pro Thr Asn Lys Met Gln Lys
385 390 395 400

Arg His Thr Phe Thr Leu Leu Val Asn Val Pro Pro Gln Ile Gly Glu

405	410	415
Asn Ala Leu Met Ala Pro Val Asp Ser Tyr Lys Tyr Gly Ser Thr Gln 420 425 430		
Ala Leu Thr Cys Thr Ile Tyr Ala Val Pro Pro Pro Ala Ala Val Leu 435 440 445		
Trp Tyr Trp Gln Leu Glu Glu Glu Cys Thr Phe Ser Pro Gln Lys Val 450 455 460		
Arg Leu Gly Ala Asn Pro Tyr Ala Cys Arg Lys Trp Lys Val Ile Ser 465 470 475 480		
Glu Arg Lys Gly Gly Asn Gln Val Glu Ile Lys Gln Arg Val Val Thr 485 490 495		
Ile Ala Gly Lys Thr Lys Thr Val Ser Thr Leu Val Ile Gln Ala Ala 500 505 510		
Asn Val Ser Ala Leu Tyr Arg Cys Met Ala Thr Asn Arg Ala Gly Ser 515 520 525		
Ser Glu Arg Val Ile Ser Phe His Val Thr Arg Gly Leu Glu Ile Asn 530 535 540		
Leu Gln Pro Arg Ser Gln Leu Thr Glu Lys Asp Asn Thr Ser Leu Gln 545 550 555 560		
Cys Thr Ala Asp Lys Phe Thr Phe Glu Lys Leu Ser Trp Tyr Lys Leu 565 570 575		
Ser Thr His Val Ser Gln Thr Pro Phe Gly Gly Leu Pro Met Pro Val 580 585 590		
Cys Lys Asn Leu Asp Ala Leu Gln Lys Leu Asn Ala Thr Val Ser Asn 595 600 605		
Val Asn Gly Glu Asn Val Thr Leu Glu Leu Ile Leu Arg Asn Ile Ser 610 615 620		
Leu Gln Asp Gly Gly Asp Tyr Val Cys Ile Ala Gln Asp Lys Lys Ala 625 630 635 640		

Lys Thr Gln His Cys Leu Val Lys His Leu Thr Val Gln Glu Pro Leu
645 650 655

His Pro Arg Leu Val Gly Asn Leu Glu Asn Gln Thr Thr Asn Ile Gly
660 665 670

Glu Thr Ile Glu Val Leu Cys Thr Val Asn Gly Val Pro Pro Pro Asn
675 680 685

Ile Thr Trp Phe Lys Asn Ser Glu Thr Leu Phe Glu Asp Ser Gly Ile
690 695 700

Val Leu Lys Asp Gly Asn Lys Thr Leu Thr Ile Arg Arg Val Arg Lys
705 710 715 720

Glu Asp Gly Gly Leu Tyr Thr Cys Leu Ala Cys Asn Ile Leu Gly Cys
725 730 735

Lys Lys Ala Glu Ala Phe Phe Ser Val Gln Gly Ala Glu Glu Lys Thr
740 745 750

Asn Leu Glu Leu Ile Ile Leu Val Gly Thr Ala Val Ile Ala Met Phe
755 760 765

Phe Trp Leu Leu Leu Val Ile Ile Leu Arg Thr Val Lys Arg Ala Asn
770 775 780

Gly Gly Asp Met Lys Thr Gly Tyr Leu Ser Ile Ile Met Asp Pro Asp
785 790 795 800

Glu Val Pro Ile Asp Glu His Cys Glu Arg Leu Pro Tyr Asp Ala Ser
805 810 815

Lys Trp Glu Phe Pro Arg Asp Arg Leu Lys Leu Gly Lys Pro Leu Gly
820 825 830

Arg Gly Ala Phe Gly Gln Val Ile Glu Ala Asp Ala Phe Gly Ile Asp
835 840 845

Lys Thr Ala Thr Cys Arg Thr Val Ala Val Lys Met Leu Lys Glu Gly
850 855 860

Ala Thr His Ser Glu His Arg Ala Leu Met Ser Glu Leu Lys Ile Leu
865 870 875 880

Ile His Ile Gly His His Leu Asn Val Val Asn Leu Leu Gly Ala Cys
885 890 895

Thr Lys Pro Gly Gly Pro Leu Met Val Ile Val Glu Tyr Cys Lys Phe
900 905 910

Gly Asn Leu Ser Ala Tyr Leu Arg Ser Lys Arg Ser Glu Phe Ile Pro
915 920 925

Tyr Lys Met Lys Ser Ala Arg Phe Arg Gln Gly Lys Glu Asn Tyr Thr
930 935 940

Gly Asp Ile Ser Thr Asp Leu Lys Gln Arg Leu Asp Ser Ile Thr Ser
945 950 955 960

Ser Gln Ser Ser Thr Ser Ser Gly Phe Val Glu Glu Arg Ser Leu Ser
965 970 975

Asp Val Glu Glu Glu Asp Ala Gly Ser Glu Asp Leu Cys Lys Asn Pro
980 985 990

Leu Thr Met Glu Asp Leu Ile Cys Tyr Ser Phe Gln Val Ala Arg Gly
995 1000 1005

Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala
1010 1015 1020

Ala Arg Asn Ile Leu Leu Ser Asp Asn Asn Val Val Lys Ile Cys
1025 1030 1035

Asp Phe Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val
1040 1045 1050

Arg Lys Gly Asp Ala Arg Leu Pro Leu Lys Trp Met Ala Pro Glu
1055 1060 1065

Thr Ile Phe Asp Arg Val Tyr Thr Ile Gln Ser Asp Val Trp Ser
1070 1075 1080

Phe	Gly	Val	Leu	Leu	Trp	Glu	Ile	Phe	Ser	Leu	Gly	Ala	Ser	Pro
1085						1090					1095			
Tyr	Pro	Gly	Val	Lys	Ile	Asp	Glu	Glu	Phe	Cys	Arg	Arg	Leu	Lys
1100						1105					1110			
Glu	Gly	Thr	Arg	Met	Arg	Ala	Pro	Asp	Tyr	Thr	Thr	Pro	Glu	Met
1115						1120					1125			
Tyr	Gln	Thr	Met	Leu	Asp	Cys	Trp	His	Gly	Asp	Pro	Lys	Gln	Arg
1130						1135					1140			
Pro	Thr	Phe	Ser	Glu	Leu	Val	Glu	His	Leu	Gly	Asn	Leu	Leu	Gln
1145						1150					1155			
Ala	Asn	Val	Arg	Gln	Asp	Gly	Lys	Asp	Tyr	Val	Val	Leu	Pro	Leu
1160						1165					1170			
Ser	Val	Ser	Leu	Asn	Met	Glu	Glu	Asp	Ser	Gly	Leu	Ser	Leu	Pro
1175						1180					1185			
Thr	Ser	Pro	Ala	Ser	Cys	Lys	Glu	Glu	Glu	Glu	Val	Cys	Asp	Pro
1190						1195					1200			
Lys	Phe	His	Tyr	Asp	Asn	Thr	Ala	Gly	Ile	Ser	Gln	Tyr	Arg	Gln
1205						1210					1215			
Gly	Ser	Lys	Arg	Lys	Ser	Arg	Pro	Val	Ser	Val	Lys	Thr	Phe	Glu
1220						1225					1230			
Asp	Ile	Pro	Leu	Val	Thr	Thr	Val	Lys	Val	Val	Gln	Glu	Glu	Asn
1235						1240					1245			
Gln	Thr	Asp	Ser	Gly	Met	Val	Leu	Ala	Ser	Glu	Glu	Leu	Lys	Thr
1250						1255					1260			
Leu	Glu	Glu	Gln	Asp	Lys	Gln	Val	Lys	Ile	Pro	Phe	Ser	Thr	Leu
1265						1270					1275			
Ala	Pro	Ser	Lys	Ser	Asn	Glu	Ser	Val	Met	Ser	Glu	Ala	Ser	Asn
1280						1285					1290			
Gln	Thr	Ser	Gly	Tyr	Gln	Ser	Gly	Tyr	His	Ser	Asp	Asp	Met	Asp

1295

1300

1305

Asn Met Val Cys Ser Ser Glu Asp Thr Glu Leu Leu Cys Ala Gln
1310 1315 1320

Glu Ala Ser Pro Thr Leu Pro Arg Cys Ala Trp Pro Gly Ile Tyr
1325 1330 1335

Ser Pro Ala Pro Val Ala Ser Leu Pro Leu
1340 1345